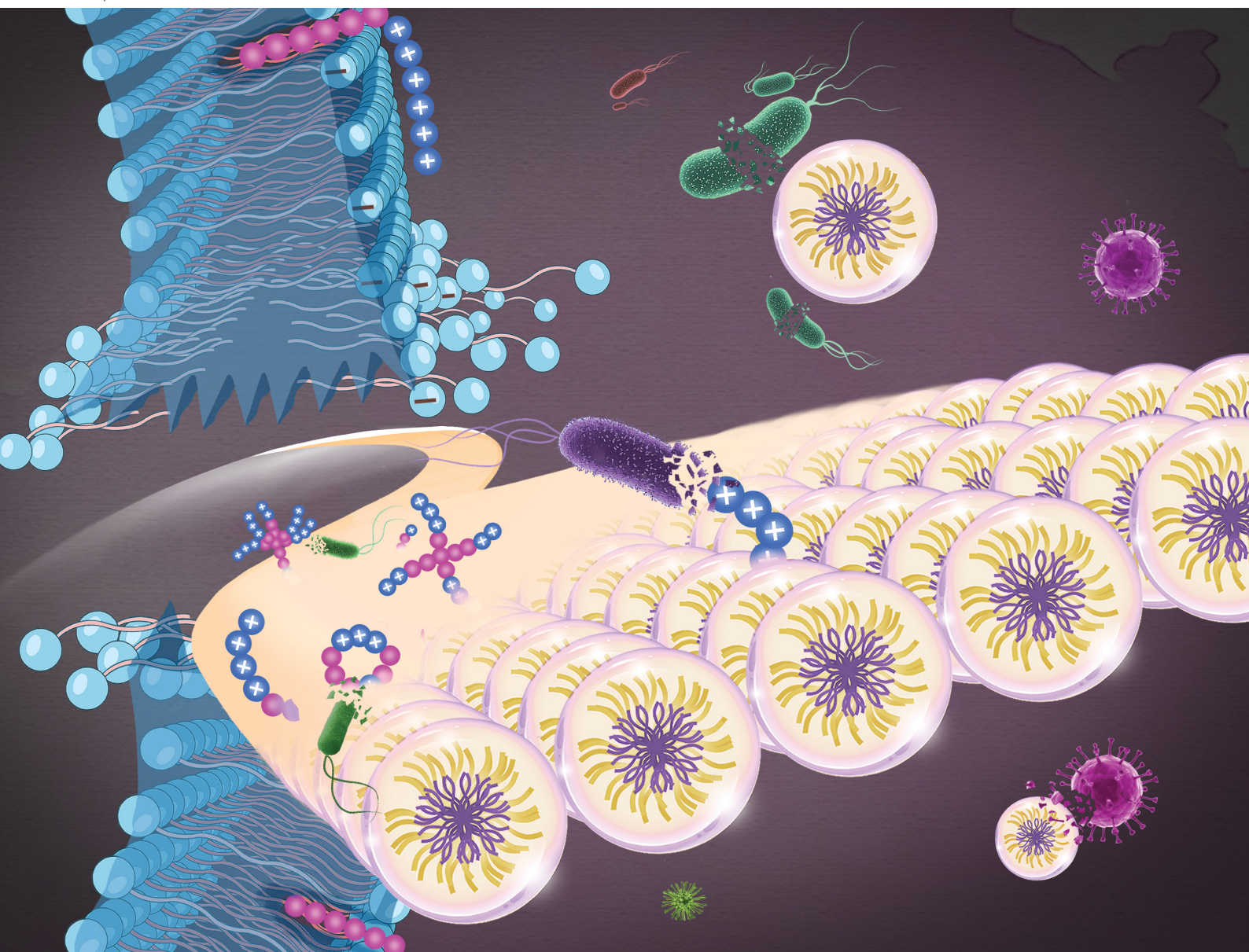


Materials Advances

rsc.li/materials-advances



ISSN 2633-5409

REVIEW ARTICLE

Zao Cheng and Patrizio Raffa
From design to application: amphiphilic copolymers as
antimicrobial materials

REVIEW

[View Article Online](#)
[View Journal](#) | [View Issue](#)

Cite this: *Mater. Adv.*, 2025,
6, 4939

Received 7th April 2025,
Accepted 5th June 2025

DOI: 10.1039/d5ma00335k

rsc.li/materials-advances

From design to application: amphiphilic copolymers as antimicrobial materials

Zao Cheng and Patrizio Raffa *

The rapid growth of harmful pathogens and their severe impact on health pose significant challenges to modern science. With comprehensive progress in human society, there is an increasing focus on personal health and quality of life, leading to a growing demand for antimicrobial materials. Amphiphilic copolymers can mimic the functions and molecular features of antimicrobial peptides (also known as host defense peptides), constituting a class of new antimicrobial materials often characterized by strongly cationic and hydrophobic domains. Due to their excellent and stable antimicrobial activity, low toxicity, and resistance prevention, amphiphilic antimicrobial copolymers have recently garnered considerable interest and attention in both academic and industrial sectors. This review outlines the latest advances of amphiphilic copolymers in antimicrobial applications, including their design strategies and current applications. Challenges and future directions of research in amphiphilic antimicrobial polymers are also discussed.

1. Introduction

As the earliest life forms on earth, microorganisms have complex and diverse metabolic mechanisms and strong adaptability to various extreme environments.^{1,2} According to statistics, many human diseases and almost all infectious ones are related to microorganisms.³ Moreover, they can also cause

corrosion and damage to various industrial equipment and raw materials, which causes huge waste of resources and economic losses.⁴ Therefore, how to inhibit the reproduction of harmful microorganisms, identify harmful bacteria, and reduce the impact of harmful microorganisms on human production and life has always been a matter of concern to humans.^{5,6} At the same time, the abuse of antibiotics has led to a significant increase in bacterial resistance, making the original antibacterial materials unable to meet people's needs.^{7–9} In recent years, researchers in related fields have been working hard to develop new materials, design various structures, and

Smart and Sustainable Polymeric Products, Engineering and Technology Institute Groningen, University of Groningen, Nijenborgh 4, 9747 AG, The Netherlands.
E-mail: p.raffa@rug.nl



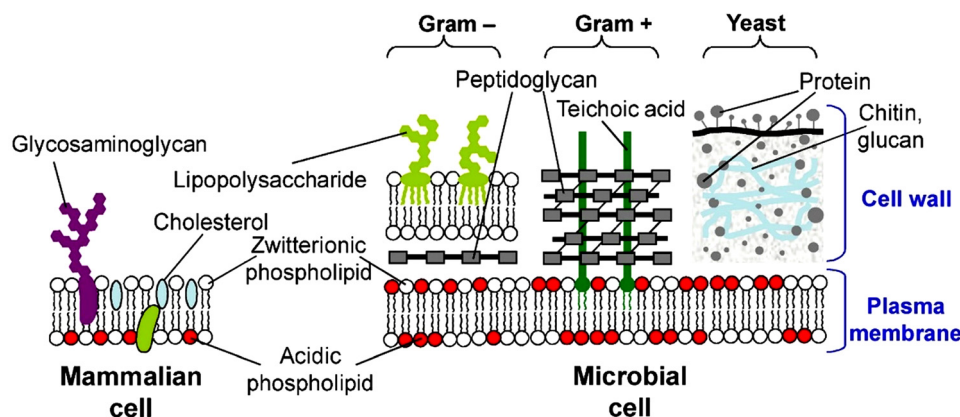
From left to right: Zao Cheng and Patrizio Raffa

November 2022. For his PhD project, he is developing novel amphiphilic polymers with antimicrobial properties, using bio-derived acrylic monomers, and controlling the polymer's architecture via RAFT polymerization. The goal of his research is to design the new generation materials for antimicrobial applications, with a particular focus on sustainability.

Patrizio Raffa (on the right) is associate professor in Smart and Sustainable Polymeric Products at the Faculty of Science and Engineering, University of Groningen, where he has been working since 2011. His current research aims at developing and synthesizing new multifunctional polymeric materials relevant for industrial and biomedical applications. In particular, he is interested in amphiphilic and stimuli-responsive polymers obtained via controlled polymerization methods, and in the use of bio-derived building blocks for the synthesis of sustainable polymeric products. His research group explores various applications for such materials as rheology modifiers, emulsifiers, dispersants, coatings, controlled release systems, pharmaceuticals & personal care products, and smart/multifunctional devices. Zao Cheng (on the left) started his PhD in Raffa's research group in

Some type of amphiphilic copolymer can mimic the functions and molecular characteristics of host defense peptides (HDPs), these typically have highly active cationic and hydrophobic chains and have shown strong antibacterial effects in many studies.^{26–28} The cationic and hydrophobic groups of these polymers are essential structures for selectively targeting and disrupting bacterial cell membranes. Achieving a fine-tuned balance in amphiphilic antimicrobial copolymers (the average quantity of hydrophobic and cationic residues present in polymer chains) is crucial for the design of highly selective antibacterial polymers.²⁹ However, These synthetic amphiphilic polymers can only mimic the primary and secondary structures of HDPs, whereas the tertiary structure of HDPs endows them with a broader range of mechanisms of action. These include inhibiting RNA synthesis,³⁰ suppressing protein synthesis,³¹ disrupting protein folding,³² interfering with lipid complexation,³³ activating microbial autolysis systems,³⁴ and neutralizing bacterial toxins.³⁵ Increasing evidence has shown that HDPs can also function by activating the host immune system.^{36,37} This involves recruiting and

Thanks to recent advancements in polymerization technologies, particularly reversible deactivation radical polymerization (RDRP), such as atom transfer radical polymerization (ATRP) and reversible addition–fragmentation chain Transfer polymerization (RAFT),^{46–49} an enormous quantity of amphiphilic antimicrobial copolymers has been developed.^{50–56} In this review, we aim to focus on the recent advances of amphiphilic copolymers for antimicrobial applications, including the design strategies for amphiphilic antimicrobial copolymers. We



© 2025 The Author(s). Published by the Royal Society of Chemistry

also aim to highlight current application directions for amphiphilic antibacterial copolymers. Finally, the challenges and development directions of amphiphilic antimicrobial copolymers are pointed out.

2. Design strategies for amphiphilic antimicrobial copolymers

At present, the research of amphiphilic antibacterial polymers focuses mainly on macromolecules obtained through the copolymerization of compounds containing cations (mostly ammonium groups) and compounds containing hydrophobic alkyl side chains; the resulting cationic amphiphilic antimicrobial copolymers have strong membrane destruction ability and have been widely used in the field of antibacterial agents.^{57–59} Many quaternary amphiphilic disinfectants have been prepared using traditional polymers, including polyvinyl pyridine,⁶⁰ polyvinyl alcohol,⁶¹ polyacrylate,⁶² and polystyrene.⁶³ Studies have shown that the bactericidal mechanism of amphiphilic antibacterial agents involves the combination of electrostatic interactions with microbial cell membranes, followed by membrane penetration caused by the insertion of hydrophobic groups. By disrupting the cell membrane, the transmembrane potential is decomposed, causing the leakage of intracellular substances and ultimately leading to cell death.^{64–67} In such models, the hydrophobicity, amphiphilic character, and the abundance of cations are considered key determinants of antibacterial activity.

This section will focus on recent researches on the impact of various components on the antibacterial activity of such amphiphilic copolymers (Fig. 2). Key aspects of localized polymer design include cationic groups,^{68,69} hydrophobic groups,⁷⁰ end groups and their amphiphilic balance.⁷¹ Additionally, the global design of polymer chains is considered, encompassing molecular weight and structural characteristics relevant to antimicrobial activity. These structural designs range from linear and cyclic structures⁷² to more advanced architectures, such as star, hyper-branched,⁷³ or brush polymers.⁷⁴ A special case of advanced architecture can be considered the sequence-

specific oligothioetheramides developed by Alabi's group, where control over the single hydrophobic units is achieved by a multi-step synthesis.^{75,76}

2.1. Local polymer design

2.1.1. Design of cationic group. The cationic functional group binds to the negatively charged bacterial cell surface through electrostatic adsorption and avoids interaction with the animal cell membrane, so it plays an irreplaceable role in this type of amphiphilic antimicrobial copolymers.⁷⁸ Therefore, in order to design efficient antibacterial polymers, many researchers began to explore the types of cationic centers and their relative spatial positions on the polymer chain, as well as the effects of their chain length and charge density on their antibacterial activity.^{51,68,71,79–83} Common polycationic structures of antimicrobial polymers include ammonium,^{45,84,85} iminium,^{82,86} phosphonium^{87,88} and sulfonium group.⁸⁹ The most common representative group of each class is discussed in this section (Table 1).

2.1.1.1. Ammonium group. The chemical structure of this class of antimicrobial agents is generally inspired by the functional and molecular characteristics of HDPs, so primary amines are the most common cationic groups, considering that lysine – with a primary amine functional group – is the main amino acid present in HDPs.⁹⁰ Several examples described here are shown in Table 1. In 2005, Kuroda⁵² reported the use of *N*-(*tert*-butoxycarbonyl)aminoethyl methacrylate and butyl methacrylate to synthesize amphiphilic random polymethacrylate derivatives with a minimum inhibitory concentration (MIC) of 16 $\mu\text{g mL}^{-1}$. In 2009, Kenichi's group prepared a library of amphiphilic random copolymers containing cationic and hydrophobic side chains by copolymerizing amine-functionalized methacrylate monomers with different proportions of alkyl methacrylates.⁶⁸ The results showed that the copolymer series with primary and tertiary amine groups exhibited strong antibacterial activity and selectivity against *Escherichia coli* (*E. coli*) versus human red blood cells (RBCs). The copolymers containing quaternary ammonium groups show activity against *E. coli* only when the hydrophobic content of the polymer is higher. c and co-workers⁹¹ reported the

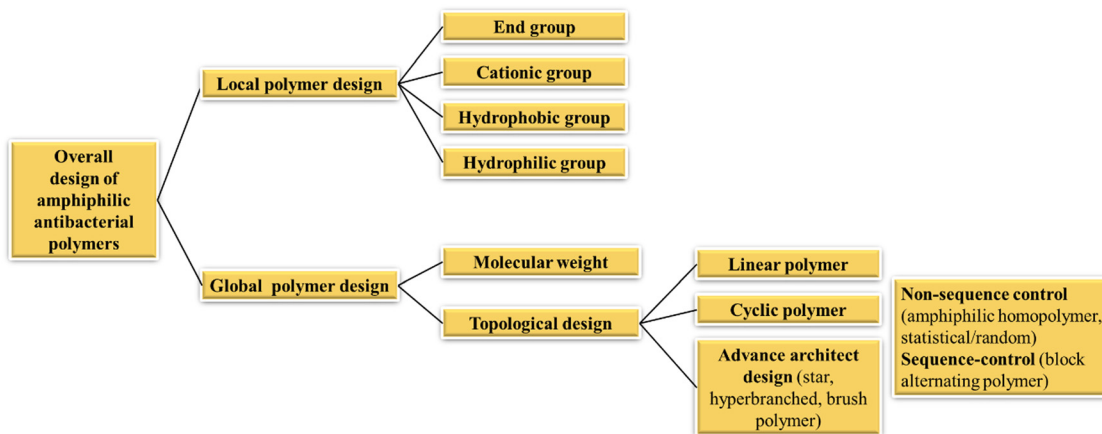


Fig. 2 Systematic design principles of amphiphilic antibacterial polymers.⁷⁷

Table 1 Chemical structures, antimicrobial activities and biocompatibility of representative amphiphilic antimicrobial polymers

Cation group	Structure	Proportions between components	Molecule weight	Antibacterial effects (MIC)	Cytotoxicity	Hemolytic for RBCs	Ref.
Ammonium		$f = 0.12^a$ $n = 8.1^a$	$M_n = 1600^a$	24 $\mu\text{g mL}^{-1}$ for <i>E. coli</i>	—	$\text{HC}_{50} = 1600 \mu\text{g mL}^{-1}$	68
		$f = 0.29^a$ $n = 12.5^a$	$M_n = 2900^a$	21 $\mu\text{g mL}^{-1}$ for <i>E. coli</i> 63 $\mu\text{g mL}^{-1}$ for <i>S. aureus</i>	—	$\text{HC}_{50} = 1300 \mu\text{g mL}^{-1}$	69
		$m = 20^c$ $n = 30$	$M_n = 9200^b$ $\text{PDI} = 1.25^b$	4.3 μM for <i>B. subtilis</i> 6.5 μM for <i>S. aureus</i> 7.0 μM for MRSA 10.8 μM for <i>E. faecalis</i> 42–63 $\mu\text{g mL}^{-1}$ for <i>S. aureus</i> 16 $\mu\text{g mL}^{-1}$ for <i>S. saprophyticus</i>	—	$\text{HC}_{50} > 54 \mu\text{M}$	91
		$n = 12^a$	$M_n = 1800^a$	—	$\text{IC}_{50} = 20 \mu\text{g mL}^{-1}$ for HEP-2 $\text{IC}_{50} = 55 \mu\text{g mL}^{-1}$ for COS-7	$\text{HC}_{50} > 1000 \mu\text{g mL}^{-1}$	92
		$x = 0.69^a$ $y = 0.31^a$ $n = 16^a$	$M_n = 2600^a$	15–60 $\mu\text{g mL}^{-1}$ for <i>S. aureus</i>	$\text{IC}_{50} > 1000 \mu\text{g mL}^{-1}$ for human fibroblasts cell	$\text{HC}_{50} > 2000 \mu\text{g mL}^{-1}$	95
Ammonium		$x = 6.7^a$ $y = 2.8^a$ $z = 7.5^a$ $n = 16.9^a$	$M_n = 2250^a$ $M_n = 3000^b$ $\text{PDI} = 1.70^b$	31 $\mu\text{g mL}^{-1}$ for <i>E. coli</i> 500 $\mu\text{g mL}^{-1}$ for <i>S. aureus</i>	—	$\text{HC}_{50} = 740 \mu\text{g mL}^{-1}$	96
		$x = 0.81^a$ $y = 0.19^a$ $n = 27^a$	4900 ^a	6.5 μM for <i>E. coli</i> 6.5 μM for <i>S. aureus</i>	$\text{IC}_{50} > 8 \text{ MIC}$ for human fibroblast cell	—	72
		$m = 41^a$ $n = 118^a$	—	—	—	—	80

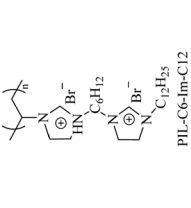
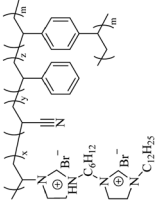
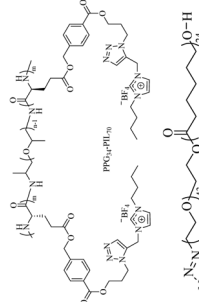
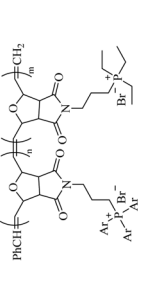
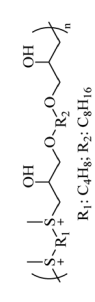
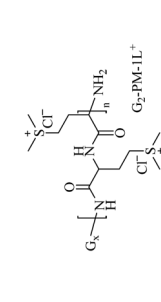




Table 1 (continued)

Cation group	Structure	Proportions between components	Molecule weight	Antibacterial effects (MIC)	Cytotoxicity	Hemolytic for RBCs	Ref.
Iminium		$x = y = 0.50^c$	$M_n = 27\,000$ $33\,000^b$	50 $\mu\text{g mL}^{-1}$ for <i>E. coli</i> 30 $\mu\text{g mL}^{-1}$ for <i>B. subtilis</i>	—	HC ₅₀ = 1245 $\mu\text{g mL}^{-1}$	102
		$x = 0.4$ $y = 0.6$	$M_n = 9326^b$	> 250 $\mu\text{g mL}^{-1}$ for <i>E. coli</i> in BM2MM 4 $\mu\text{g mL}^{-1}$ for <i>P. aeruginosa</i> in BM2MM	—	HC ₅₀ = 551.6 $\mu\text{g mL}^{-1}$	106
		$x = 0.79^a$ $y = 0.21^a$	$M_n = 20\,400^a$ $M_n = 13\,000^b$ PDI = 1.14 ^b	312 $\mu\text{g mL}^{-1}$ <i>S. aureus</i> 5000 $\mu\text{g mL}^{-1}$ <i>E. coli</i> 312 $\mu\text{g mL}^{-1}$ <i>S. epidermidis</i> 2500 $\mu\text{g mL}^{-1}$ <i>P. aeruginosa</i> 312 $\mu\text{g mL}^{-1}$ <i>C. parapsilosis</i>	—	HC ₅₀ = 2500 $\mu\text{g mL}^{-1}$	104
		$n = 20^c$	—	7.8 $\mu\text{g mL}^{-1}$ for <i>E. coli</i> 7.8 $\mu\text{g mL}^{-1}$ for <i>K. pneumoniae</i> 15.6 $\mu\text{g mL}^{-1}$ for <i>P. aeruginosa</i> 15.6 $\mu\text{g mL}^{-1}$ for <i>C. albicans</i> 3.9 $\mu\text{g mL}^{-1}$ for <i>S. aureus</i>	—	—	100
Iminium		$x = 6$; molecular weight of $n = 7000^c$ and $m = 3000^c$	$M_n = 11\,459^a$ $M_n = 5630^b$ PDI = 3.24 ^b	256 $\mu\text{g mL}^{-1}$ for <i>E. coli</i> 64 $\mu\text{g mL}^{-1}$ for <i>S. aureus</i>	—	HC ₅₀ > 1000 $\mu\text{g mL}^{-1}$	103
		—	$M_n = 6500^b$	25 $\mu\text{g mL}^{-1}$ for <i>E. coli</i> 50 $\mu\text{g mL}^{-1}$ for MRSA	—	HC ₅₀ = 740 $\mu\text{g mL}^{-1}$	105

Table 1 (continued)

Cation group	Structure	Proportions between components	Molecule weight	Antibacterial effects (MIC)	Cytotoxicity	Hemolytic for RBCs	Ref.
Iminium		—	$M_n = 12\,000^b$ $PDI = 1.02^b$	$0.018\ \mu\text{g mL}^{-1}$ for <i>E. coli</i> $0.009\ \mu\text{g mL}^{-1}$ for <i>S. aureus</i>	—	—	107
		—	—	The antibacterial rate against <i>E. coli</i> and <i>S. aureus</i> > 90%	—	Hemolytic activity (<4%)	107
		—	$M_n = 26\,300^b$ $PDI = 1.18^b$	$25\ \mu\text{g mL}^{-1}$ for <i>E. coli</i> $25\ \mu\text{g mL}^{-1}$ for <i>S. aureus</i>	$IC_{50} > 100\ \mu\text{g mL}^{-1}$ for NIH 3T3 $IC_{50} > 100\ \mu\text{g mL}^{-1}$ for 293T	Hemolytic activity <5% at $100\ \mu\text{g mL}^{-1}$	108
Phosphonium		—	$M_n = 6500^b$ $PDI = 1.12^b$	$0.13\ \text{mM}$ (MBC) for <i>S. aureus</i> and <i>E. coli</i>	—	NO hemolytic at $0.53\ \text{mM}$ (sheep red blood cells)	110
Phosphonium		Phenyl:ethyl (5:5) $m/n = 0.73^d$	$M_n = 13\,131^a$ $M_n = 6590^b$ $PDI = 3.59^b$	$128\ \mu\text{g mL}^{-1}$ for <i>E. coli</i> $32\ \mu\text{g mL}^{-1}$ for <i>S. aureus</i>	—	$HC_{50} = 2048\ \mu\text{g mL}^{-1}$	111
Sulfonium		—	$M_n = 10\,500^b$ $PDI = 1.53^b$	$1.25\ \mu\text{g mL}^{-1}$ (MBC) for <i>S. aureus</i> $1.25\ \mu\text{g mL}^{-1}$ (MBC) for MRSA $10\ \mu\text{g mL}^{-1}$ (MBC) for <i>E. faecalis</i> $2.5\ \mu\text{g mL}^{-1}$ (MBC) for <i>E. coli</i> $20\ \mu\text{g mL}^{-1}$ (MBC) for <i>P. aeruginosa</i>	—	$HC_{10} = 600\ \mu\text{g mL}^{-1}$	87
		$x = 2$ $n = 39^b$	$M_n = 94\,410^b$ $PDI = 1.37^b$	$8\ \mu\text{g mL}^{-1}$ for <i>E. coli</i> $16\ \mu\text{g mL}^{-1}$ for <i>S. aureus</i>	—	$HC_{10} = 600\ 32\,000\ \mu\text{g mL}^{-1}$	79

G_x: x=0, 1, and 2, representing 4, 8, 16 arms

Table 1 (continued)

Cation group	Structure	Proportions between components	Molecule weight	Antibacterial effects (MIC)	Cytotoxicity	Hemolytic for RBCs	Ref.
		$n = 28^b$	$M_n = 1270^b$ $PDI = 1.6^b$	$10 \mu\text{g mL}^{-1}$ for <i>E. coli</i> $5 \mu\text{g mL}^{-1}$ for <i>S. aureus</i> $10 \mu\text{g mL}^{-1}$ for <i>M. smegmatis</i>	—	$\text{HC}_{50} = 200 \mu\text{g mL}^{-1}$ (sheep red blood cells)	88

RBCs: human red blood cells; IC₅₀: the maximum concentration of polymer less than 50% toxic to test cells; HC₅₀ and HC₁₀: the maximum concentration of polymers with less than 50% or 10% hemolytic activity on red blood cells; The polymer characterization data listed in the table are the polymers with the highest antibacterial activity in the corresponding article. ^a The values were calculated from analysis of the peak integration of the ¹H NMR spectra. ^b The values were determined by GPC. ^c The values were theoretical.

Quaternary ammonium has permanent cationic charges, regardless of the pH, and they are also commonly used in such amphiphilic antimicrobial polymers.⁹⁷ Hedrick and Yang *et al.*⁹⁸ reported the preparation of a series of antibacterial polyionenes by utilizing cations brought by two-sided quaternary amines, such as

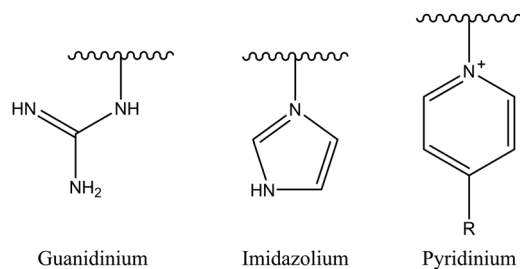


Fig. 3 Schematic diagram of the iminium group structure.

bis[2-(*N,N*-dimethylamino)ethyl] ether, through a simple addition polymerization reaction, in which the polymer formation reaction and charge installation occurred simultaneously. Tyagi and Mishra⁷² used aminopropyl methacrylamide (APMA), which remains protonated at physiological pH, thus providing a cationic charge, while benzylmethacrylamide (BMA) was chosen to provide hydrophobicity due to the presence of an aromatic hydrophobic benzyl group. Wang *et al.*⁸⁰ synthesized a series of quaternized triblock copolymers that can self-assemble into reverse micelles (RM) in tetrahydrofuran (THF). The outer shell of the RM contains biocompatible poly(ϵ -caprolactone) (PCL) blocks to provide bio-safety and responsiveness, and the core contains quaternary ammonium cationic blocks to provide antimicrobial activity. In the presence of bacterial lipase, the biodegradable PCL blocks are hydrolyzed, resulting in the responsive release of quaternary biocidal agents (QBA), thereby achieving self-sterilization. This provides a promising strategy for the development of bio-switchable self-sterilizing dressing materials.

2.1.1.2. Iminium group. Another example of nitrogen-based cationic center is the iminium group. This classification comprises several candidates, such as pyridinium, imidazolium, and guanidinium salts (Fig. 3). The iminium group exhibits uniformly distributed positive charges *via* π bonds and aromatic conjugated systems, potentially augmenting its affinity for negatively charged bacterial membranes and consequently enhancing antibacterial effectiveness.²⁹

Among iminium groups, guanidinium salts are most widely used as cationic charge sources for such amphiphilic antimicrobial copolymers because of their close structural similarity to the natural amino acid arginine. Some studies have found that guanidine-functionalized polycarbonate can pass through bacterial cell membranes without damaging their integrity, and non-specifically binds to intracellular proteins and nucleic acids.⁹⁹ Therefore, Leong *et al.* simply mixed quaternary ammonium homopolymer and guanidinium homopolymer to produce a more effective antibacterial behavior than a single mechanism, showing a synergistic or additive effect.¹⁰⁰ An increased antibacterial activity for guanidine-containing oligothioetheramides, compared to other ammonium groups, was reported by Alabi's research group.¹⁰¹

Using a pyridine ring as a source of cations, Sambhy *et al.*¹⁰² investigated the effect of the spatial relationship of the charges and alkyl tails on the polymer backbone on membrane disruptive capacity. The results showed that placing the charges

and tails in a spatially separated center resulted in higher membrane disruptive capacity, as evidenced by enhanced antimicrobial and hemolytic activities. Islam *et al.*¹⁰³ synthesized random and block dendritic oxanorbornene polymers with different hydrophobicity/hydrophilicity and with pyridinium salt groups using ring-opening metathesis polymerization (ROMP), in which the copolymers with a six-carbon linker, a high charge density, and hexylpyridine functional groups per repeating unit, exhibited high antibacterial activity against Gram-positive bacteria (*S. aureus*). However, all polymers were inactive against Gram-negative bacteria (*E. coli*). Cuervo-Rodríguez *et al.*¹⁰⁴ quaternized polymers containing thiazole ring monomers with butyl iodide or octyl iodide to obtain cationic amphiphilic antimicrobial copolymers containing thiazolium groups, and utilized the different alkyl groups chains length to adjust their hydrophobic/hydrophilic balance. The results showed that the more hydrophobic octanoylated copolymer was more effective against all tested microorganisms. Hancock *et al.*¹⁰⁵ synthesized *N*-methylpyridinium-fused norbornene monomers through ROMP, and successfully synthesized backbone cationic polymers with diverse structures, showing that these polymers are effective against Gram-negative (*E. coli*) and Gram-positive (*S. aureus*) bacteria, with MIC as low as 25 $\mu\text{g mL}^{-1}$. In addition, polymers with a smaller DP showed increased selectivity for bacteria over human RBCs. Vishwakarma *et al.*¹⁰⁶ reported the development of water-soluble synthetic peptide-mimicking polyurethanes incorporating arginine as a cationic functional group. The results demonstrated their ability to disrupt pre-formed biofilms of *P. aeruginosa*, *S. aureus*, and *E. coli*. Although these polyurethanes exhibited relatively weak antibacterial activity against planktonic bacteria, they effectively prevented surface adhesion and promoted bacterial surface motility. This dual action inhibited biofilm formation of both Gram-positive and Gram-negative bacteria at sub-inhibitory concentrations, while showing no toxicity toward mammalian cells.

Using imidazole groups as cation sources, Zheng *et al.*¹⁰⁷ synthesized imidazolium-type ionic liquid monomers and their corresponding poly(ionic liquids) (PILs) membranes. By studying the impact of chemical on antibacterial activity, the results show that the antibacterial activity increases with the increase in charge density (double cation), while decreases with increasing carbon chain length. Shi *et al.*¹⁰⁸ using ultrafast ring-opening polymerization (ROP) technology and "click" chemical reactions, prepared imidazolium-based block co-polypeptides with L configuration (PPGn-PILm); these can self-assemble into positively charged nanoscale micelles, and showed MIC against both Gram-positive (*S. aureus*) and Gram-negative (*E. coli*) of 25 $\mu\text{g mL}^{-1}$.

2.1.1.3. Phosphonium group. Since the structure of phosphonium compounds is similar to that of quaternary ammonium ones, Kanazawa *et al.*¹⁰⁹ first reported the use of phosphonium as a cationic charge source for amphiphilic antimicrobial copolymers in 1993. It was also shown that for polymers with the same structure but different cationic parts, the antibacterial activity of phosphates is two orders of magnitude higher than that of polymer ammonium salts. Hisey *et al.*¹¹⁰ connected



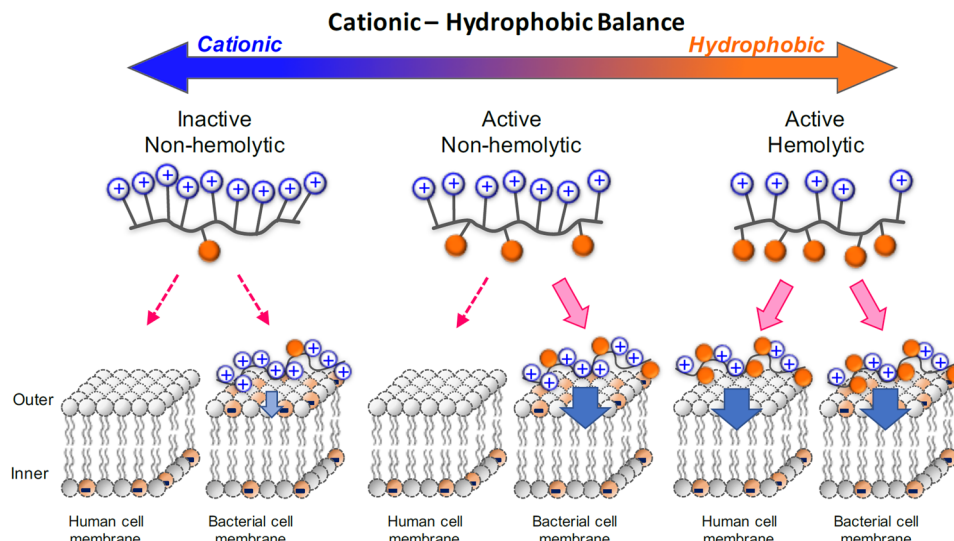


Fig. 4 Schematic diagram illustrating the impact of cationic-hydrophobic balance on the cell selectivity of amphiphilic antimicrobial copolymers. Reproduced with permission from ref. 81, copyright 2017, American Chemical Society.

phosphonium cations with different alkyl lengths to the ends of polyethylene glycol-polycaprolactone block copolymers and allowed the phosphonium-functionalized block copolymers to self-assemble in aqueous solution forming micro micelles to achieve selective antibacterial function. Kudah *et al.*¹¹¹ synthesized five different aromatic side-chain phosphonium-based polymers *via* cyclopentene ROMP; the synthesized polymers showed higher antibacterial activity against Gram-positive strains than Gram-negative strains.

2.1.1.4. Sulfonium group. In 1993, Kanazawa *et al.*¹¹² synthesized polymers with antibacterial activity using sulfur atoms as cation sources. In recent years, sulfonium compounds have attracted attention as sources of cationic charge for amphiphilic antimicrobial copolymers because they are similar in charge to quaternary ammonium groups and are also commonly used as therapeutic agents.^{79,87–89} Research results have shown that when the polymers contain sulfonium as a cationic moiety in the backbone, they can exhibit enhanced antimicrobial activity and selectivity compared to corresponding polymers with sulfonium as side chains. This makes the former potential candidates for combating antibiotic-resistant bacterial infections.^{87,88}

Hu *et al.*⁸⁷ developed a new class of AB alternating sequence main chain sulfonium-containing polymers. Broad-spectrum bactericidal properties against a group of clinically relevant bacteria and excellent biocompatibility were observed for the first time in a sulfonium-based macromolecule. This amphiphilic polymer showed excellent antibacterial activity against clinically relevant bacteria, including MRSA, with MIC in the range of 1.25–10 $\mu\text{g mL}^{-1}$ and no hemolysis even at polymer concentrations as high as 10 000 $\mu\text{g mL}^{-1}$. Oh and Khan⁸⁸ synthesized β -hydroxy-functionalized sulfonium polymers, which demonstrated structural stability and high efficacy against clinically significant Gram-positive bacterial strains, such as *Mycobacterium smegmatis*, which is crucial for

combating tuberculosis infections. Furthermore, they are equally effective against challenging Gram-negative bacterial strains, such as *E. coli*. Additionally, the developed main-chain sulfonium polymers show similar efficacy to antibiotics like kanamycin and vancomycin.

In addition, Zhang *et al.*⁷⁹ synthesized a series of star-shaped alkylated sulfonium amphiphilic antimicrobial copolymers by mimicking the structure unit of vitamin U *via* *N*-carboxyanhydrides (NCA) ring-opening reactions. The short hydrophobic moiety and highly cationic structure significantly enhance the hemocompatibility and bactericidal activity of G₂-PM-1^{H+} (Table 1, HC₁₀/MIC ratio as high as 16 000). G₂-PM-1^{H+} has broad-spectrum bactericidal activity (*S. aureus*: MIC = 4 $\mu\text{g mL}^{-1}$, *E. coli*: MIC = 16 $\mu\text{g mL}^{-1}$) and has excellent drug resistance compared with vancomycin.

In general, the full potential of sulfonium compounds as sources of cationic charge is still to be explored, especially sulfonium-based polymers with well-defined sequences.

2.1.2. Hydrophobic group. As the active components directly involved in the insertion and disruption of bacterial and mammalian membranes, the local hydrophobicity of monomers and the overall hydrophobicity of polymers (*i.e.*, global amphiphilic balance) represent characteristic parameters reflecting the influence of hydrophobic groups on biological activity.^{45,53,71,113} Therefore, the balance between hydrophilicity and hydrophobicity is one of the key determinants for amphiphilic antimicrobial copolymers (Fig. 4). This balance can be optimized at the local monomer level and/or the overall polymer level in amphiphilic antimicrobial copolymers or homopolymers.^{114–116} To achieve this, designers need to consider the hydrophobic contribution of the polymer backbone as well as the characteristics of the side chains (such as hybridization, branching, and carbon number). Three primary strategies can be employed to adjust this balance, namely: (1) managing the structure of monomers, such as the length



of spacer units and the type of hydrophobic groups; (2) adjusting the hydrophobicity ratio relative to other functional groups in amphiphilic antimicrobial copolymers; (3) inserting neutral hydrophilic groups, such as poly ethylene glycol (PEG).

The first design strategy involves controlling the hydrophobicity at the local level of the polymer by designing the structure of different hydrophobic monomers, thereby regulating the hydrophilic–hydrophobic balance of the polymer. So far, most studies on antimicrobial polymers have utilized hydrophobic alkyl chains as their hydrophobic groups. Therefore, the structural properties of the hydrophobic alkyl tails, such as the length of alkyl chains and the type of side chains, are crucial for determining the biological activity of the polymer. According to reports,^{56,81,86,104,113,114,117–119} extending the chain length within a certain range of carbon atoms in the side chains results in increased antibacterial activity. Hydrophobic structures can interact more strongly with the lipid bilayer of microbial membranes, thereby enhancing antibacterial activity. However, excessively long alkyl chains may lead to polymer over-hydrophobicity, potentially causing reduced water solubility and increased indiscriminate toxicity to pathogens and host cells, *i.e.*, enhanced cytotoxicity. Therefore, optimizing the length of hydrophobic chains helps to regulate the balance between antibacterial efficacy and cytotoxicity, thereby enhancing the bioavailability of amphiphilic antimicrobial copolymers and maximizing their antibacterial activity.

In addition to hydrophobic chain length, the type of hydrophobic part also influences the bioactivity.^{50,113} Tan *et al.*¹²⁰ synthesized a series of guanidinium-functionalized random polycarbonates through organocatalytic ring-opening polymerization (Fig. 5).

By introducing various length of hydrophobic co-monomers (ethyl, propyl, isopropyl, benzyl, and hexyl) at different percentage contents, they modulated the hydrophobicity of the polymers. Overall, these polymers exhibited similar minimum inhibitory concentrations and minimum bactericidal concentrations against a broad spectrum of microbes. However, polymers with higher hydrophobicity showed a faster rate of bacterial elimination. At higher percentages (20 mol%), polymers containing a higher proportion of hydrophobic co-

monomers displayed reduced selectivity due to their elevated hemolytic activity (Table 2).

A series of antimicrobial surfactant-like poly(ester polyurthane)s with different hydrophobic groups (short alkyl chain (PSA), aromatic group (PA) and long alkyl chain (PLA)) and cationic groups uniformly distributed along the polymer chain were studied (Fig. 6).¹²² Membrane permeability assays showed that all three antimicrobial polymers were able to permeabilize the outer (periplasm) and inner membrane (cytoplasm) of *E. coli* and that the *S. aureus* cell wall was unable to prevent these polymers from damaging the cytoplasmic membrane. PA and PLA caused significant damage to the plasma membrane of *E. coli*, whereas PSA caused slight damage to the integrity of the plasma membrane, but even this level of damage was able to dissipate the plasma membrane potential and lead to cell death (Table 3).

The second and third design approaches focus more on the global polymer level, wherein the hydrophobic group influences bioactivity through its interaction with other comonomers present in the global amphiphilic balance of amphiphilic antimicrobial copolymers.

For example, Phuong *et al.*¹¹³ synthesized a library of 36 amphiphilic antimicrobial polymers using eight hydrophobic monomers (cyclic, aromatic, linear, or branched and classified into three categories, C5, C7, and C9, based on the carbon number (5, 7, or 9)) (Fig. 7) and screened them against four bacterial strains and sheep RBCs to assess their antimicrobial activity and blood compatibility. For polymers with saturated chains and similar DP and target component molar ratios, the antimicrobial activity decreases as the carbon number of the hydrophobic side chain groups increases (*i.e.*, the MIC value for the C9 group is higher than those for C7 and C5). Within the same polymer group, copolymers containing branched monomers exhibit better antimicrobial activity than those with linear monomers. In the C5 group (see Fig. 7), the length of **Im** (6.2 Å) is slightly shorter than that of **Pm** (7.4 Å), and the MIC value of I40-1535 against Gram-negative bacteria is slightly lower than that of P40-1535. In the C7 group, the difference in chain length between branched (**Pbm**, 5 Å) and linear (**Hm**, 9.9 Å) monomers is significant, resulting in a substantial difference in

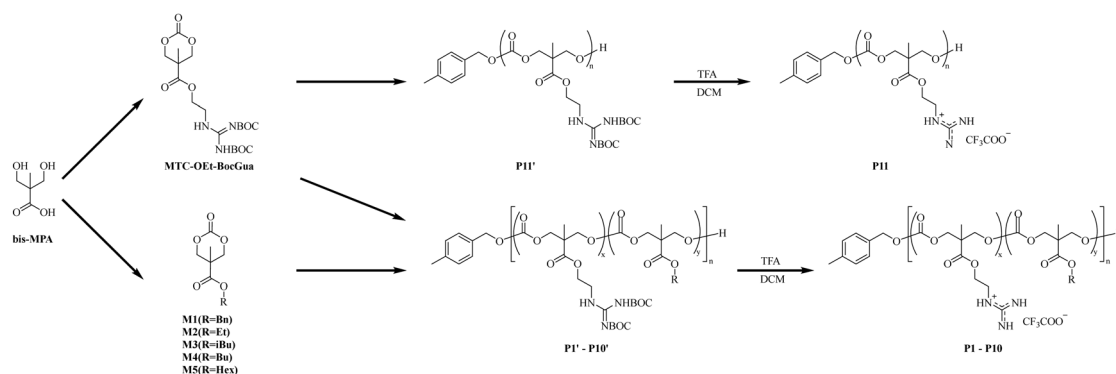
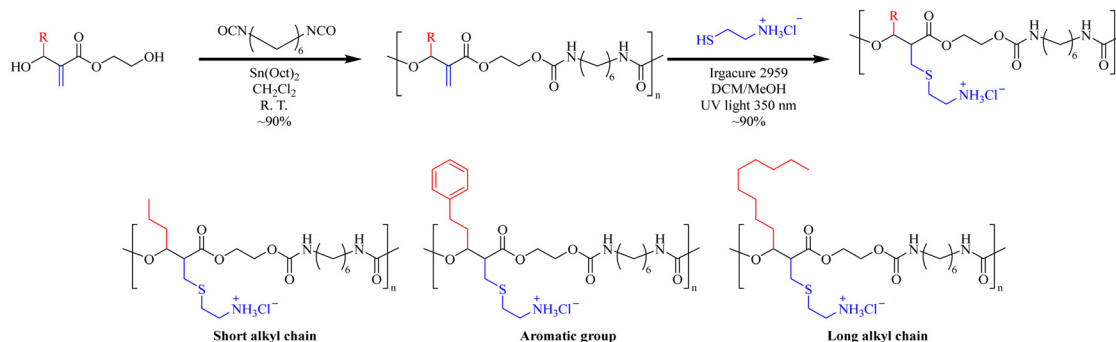


Fig. 5 Synthesis of guanidinium-functionalized random polycarbonates with different hydrophobic side chain lengths via organocatalytic ring-opening polymerization. Redrawn from ref. 120.

Table 2 Minimum inhibitory concentrations (MIC, $\mu\text{g mL}^{-1}$) and selectivity of the random polycarbonates with different hydrophobic side chain lengths against a broad spectrum of microbes. Redrawn from ref. 120

Polymer	R	x	y	MIC [$\mu\text{g mL}^{-1}$]/selectivity (HC ₅₀ /MIC)						HC ₅₀ [$\mu\text{g mL}^{-1}$]
				Gram-positive		Gram-negative			Fungus	
				<i>S. aureus</i>	<i>A. baumannii</i>	<i>E. coli</i>	<i>K. pneumoniae</i>	<i>P. aeruginosa</i>	<i>C. albicans</i>	
P1	Bn	17	2	7.8 (>256)	7.8 (>256)	7.8 (>256)	15.6 (>128)	15.6 (>128)	31.3 (>64)	>2000
P2	Et	18	2	7.8 (>256)	7.8 (>256)	7.8 (>256)	15.6 (>128)	15.6 (>128)	31.3 (>64)	>2000
P3	iBu	16	2	7.8 (>256)	7.8 (>256)	7.8 (>256)	15.6 (>128)	15.6 (>128)	31.3 (>64)	>2000
P4	Bu	18	2	7.8 (>256)	7.8 (>256)	7.8 (>256)	15.6 (>128)	15.6 (>128)	31.3 (>64)	>2000
P5	Hex	18	2	7.8 (>256)	7.8 (>256)	7.8 (>256)	15.6 (>128)	15.6 (>128)	31.3 (>64)	>2000 ^a
P6	Bn	15	4	7.8 (51)	7.8 (51)	7.8 (51)	15.6 (26)	15.6 (26)	31.3 (13)	≈400
P7	Et	15	4	7.8 (>256)	7.8 (>256)	7.8 (>256)	15.6 (>128)	15.6 (>128)	31.3 (>64)	>2000
P8	iBu	14	4	7.8 (51)	7.8 (51)	7.8 (51)	15.6 (29)	15.6 (29)	31.3 (14)	≈450
P9	Bu	13	5	7.8 (>256)	7.8 (>256)	7.8 (>256)	15.6 (>128)	15.6 (>128)	31.3 (>64)	>2000 ^a
P10	Hex	14	4	7.8 (22)	15.6 (11)	7.8 (22)	15.6 (11)	15.6 (11)	31.3 (5)	≈170
P11	—	18	—	15.6 (>513)	7.8 (>1026)	15.6 (>513)	31.3 (>256)	15.6 (>513)	31.3 (>256)	>8000 ¹²¹

^a At 2000 $\mu\text{g mL}^{-1}$, the hemolytic activity of P5 and P9 is ≈45% and ≈37%, respectively.

**Fig. 6** Synthesis of antimicrobial surfactant-like poly(ester polyurethane)s with different hydrophobic pendant groups (PSA, PA, and PLA) and cationic groups distributed uniformly along the polymer chain. Redrawn from ref. 122.**Table 3** MIC of the Poly(ester polyurethane)s with different hydrophobic pendant groups (PSA, PA, and PLA) against Gram-negative/positive bacteria and HC₁₀ toward RBCs. Redrawn from ref. 122

Polymer	MIC									
	Gram-negative bacteria					Gram-positive bacteria				HC ₁₀
	<i>E. coli</i>		<i>P. aeruginosa</i>		<i>S. aureus</i>		<i>S. epidermidis</i>			
	µg mL ⁻¹	µM	µg mL ⁻¹	µM	µg mL ⁻¹	µM	µg mL ⁻¹	µM		
PSA	16	2.0	32	4.0	32	4.0	16	2.0	379.4	47.4
PA	8	1.6	16	3.1	16	3.1	4	0.8	19.8	3.8
PLA	8	0.9	16	1.8	16	1.8	16	1.8	3.5	0.4
Ampicillin	1	2.9	128	371.2	0.2	0.6	4	11.6	—	—
Melittin	25	8.8	50	17.6	25	8.8	0.8	0.3	—	—

antimicrobial activity (Table 4). **Im** monomer shows the highest compatibility with the selected cationic and hydrophilic monomers, as evidenced by polymers I40-1535 and I40-1040, which exhibit the highest antimicrobial activity among the tested polymers. In summary, it was found that linear and branched alkyl hydrophobic monomers exhibited better antimicrobial effects than cyclic and aromatic ones, and polymers with a Calculation $\log P$ (the partition constant of a compound between *n*-octanol (hydrophobic phase) and water (hydrophilic

phase)) between 0 and 2 were most likely to achieve the optimal balance of high antimicrobial activity and low hemolytic activity.

Another notable example comes from Mankoci *et al.*,¹²³ which reported a series of water-soluble antimicrobial polyurethanes designed to mimic antimicrobial peptides (AMPs) (Fig. 8). These polyurethanes were synthesized by incorporating cationic side chains into one segment of the polymer chain, while functionalizing the other segment with various side-chain



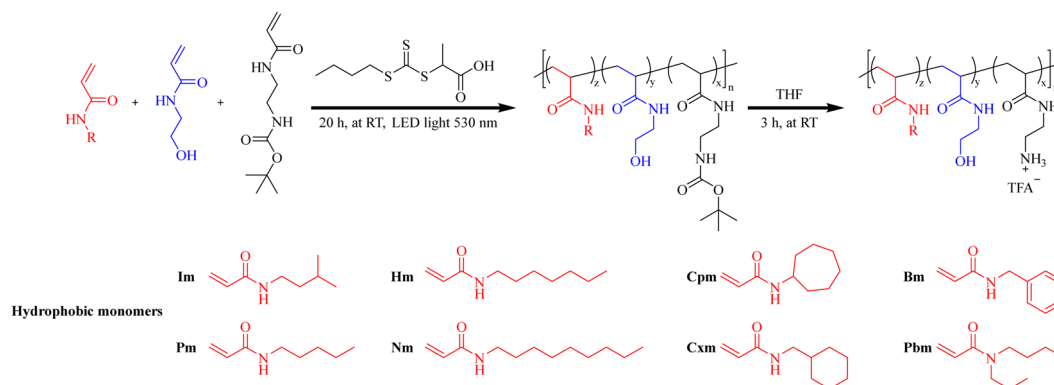


Fig. 7 Synthesis of amphiphilic antimicrobial polymers using eight hydrophobic monomers (*N*-isopentylacrylamide (**Im**), *N*-pentylacrylamide (**Pm**), *N*-heptylacrylamide (**Hm**), *N*-butyl-*N*-propylacrylamide (**Pbm**), *N*-cycloheptylacrylamide (**Cpm**), *N*-(cyclohexylmethyl)acrylamide (**Cxm**), *N*-benzylacrylamide (**Bm**), and *N*-nonylacrylamide (**Nm**)). Redrawn from ref. 113.

Table 4 Antimicrobial activities (MIC) of 40-DP_n polymers with different hydrophobic monomers and monomer feed ratios against four microorganisms. Redrawn from ref. 113

Family of polymers	Hydrophobic monomer (<i>C log P</i>)	Polymer	Monomer ratio (<i>x:y:z</i>)	ClogP of representative oligomer	MIC of polymer (μg mL ⁻¹)			
					<i>PA01</i>	<i>P. aeruginosa</i>	<i>E. coli</i>	<i>S. aureus</i>
I-family	Im (1.40)	I40-1040	50:10:40	2.70	16	16–32	16	>256
		I40-1535	50:15:35	1.55	16	16–32	16	>256
		I40-2030	50:20:30	0.39	16–32	32–64	32	>256
		I40-2525	50:25:25	−0.77	256	>256	128–256	>256
		I40-3020	50:30:20	−1.93	>256	>256	>256	>256
P-family	Pm (1.53)	P40-1040	50:10:40	3.22	32	32	16	>256
		P40-1535	50:15:35	2.00	32	32	16	>256
		P40-2030	50:20:30	0.78	32	32–64	32–64	>256
H-family	Hm (2.59)	H40-2030	50:20:30	3.95	64	128	64	>256
		H40-2525	50:25:25	2.2	64	64	64	>256
		H40-3020	50:30:20	0.45	64–128	128–256	128	>256
Pb-family	Pbm (2.37)	Pb40-2030	50:20:30	3.70	32	32–64	16–32	>256
		Pb40-2525	50:25:25	2.00	32–64	32–64	32–64	>256
		Pb40-3020	50:30:20	0.29	64	128	128–256	>256
Cp-family	Cpm (2.01)	Cp40-2030	50:20:30	2.20	32–64	64	32	>256
		Cp40-2525	50:25:25	0.74	32–64	32–64	16–32	>256
Cx-family	Cxm (2.07)	Cx40-2030	50:20:30	2.38	32	32–64	32	>256
		Cx40-2525	50:25:25	0.89	32–64	64	16–32	>256
B-family	Bm (1.18)	B40-1535	50:15:35	1.11	32–64	64	64–128	>256
		B40-1040	50:10:40	2.23	16–32	32	32	>256
		B40-2030	50:20:30	0.05	64	64–128	256	>256
N-family	Nm (3.65)	N40-2030	50:20:30	7.13	>256	>256	>256	>256
		N40-3020	50:30:20	2.57	128–256	256	128–256	>256
		N40-3515	50:35:15	0.29	128–256	256	128–256	>256

amino acids. Their antimicrobial activity and cytotoxicity were systematically compared. The findings revealed that the addition of hydrophobic, neutral polar, or anionic components did not significantly influence antibacterial activity. However, the introduction of hydrophobic groups enhanced cytoplasmic membrane disruption, and the proportion of cationic side-chain groups correlated with the degree of outer membrane disruption in Gram-negative bacteria. Meanwhile, neutral polar and anionic groups improved the compatibility of the polyurethanes with mammalian cells, highlighting the tunability of these materials for selective activity and reduced cytotoxicity.

Typical HDPs feature two functionalities, cationic groups and hydrophobic moieties. However, as mentioned above, an

excess of cationic iminium^{82,86–88} and hydrophobic^{113,120,122} characteristics can lead to indiscriminate toxicity towards all cell types, including RBCs. This renders the optimization of binary polymer balance challenging. Consequently, a recent approach has been the introduction of neutral hydrophilic groups, resulting in a class of ternary polymers. The presence of neutral hydrophilic groups serves to reduce undesired protein complexation and hemolysis, thus maintaining polymer antimicrobial activity while imparting biocompatibility (Fig. 9).

The role of the neutral hydrophilic group was reported first by Allison *et al.*¹²⁵ They focus on quaternized poly(vinyl-pyridine) (PVP), which inherently possesses antibacterial properties but also exhibits high toxicity. They found that incorporation of



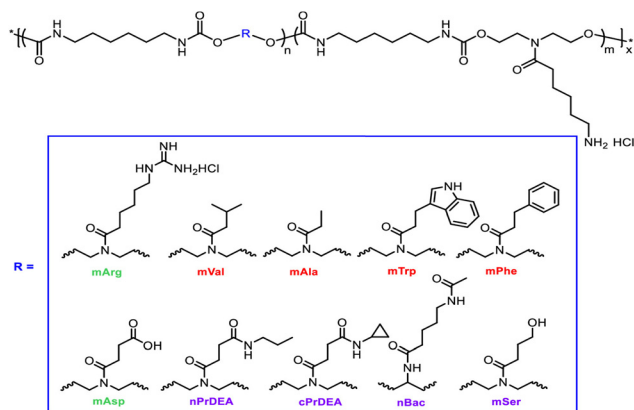


Fig. 8 Chemical structures of the antimicrobial polyurethanes. **mArg** and **mAsp** (in green) represent the charged polar pendant groups; **mVal**, **mAla**, **mTrp**, and **mPhe** (in red) represent the hydrophobic pendant groups, and **nPrDEA**, **cPrDEA**, **nBac**, and **mSer** (in purple) were chosen as uncharged polar pendant groups for antimicrobial polyurethane design. Redrawn from ref. 123.

polyethylene glycol comonomers could reduce hemolytic activity without compromising antimicrobial efficacy. This study paves the way for the development of new ternary polymer systems akin to HDPs. Since then, PEG has been commonly employed as a neutral hydrophilic residue.^{70,126–128} Additionally, various other neutral hydrophilic candidates exist for ternary polymer systems, such as hydroxyl substitutions (mimicking serine residues).^{45,129} Based on previous research, Pham *et al.*⁷⁰ designed and developed a series of statistical amphiphilic ternary copolymers and systematically investigated the effects of each component group on antimicrobial activity and biocompatibility (Fig. 10).

The results indicate that, unlike hydrophobic moieties that directly damage cell membranes, hydrophilic groups indirectly but significantly influence bioactivity by modulating the hydrophobic/hydrophilic balance and overall hydrophobicity, leading to alterations in polymer solubility properties. The AM family (Fig. 10) exhibited the highest antibacterial activity against Gram-negative bacteria. The **PEG-A** and **PEG-AA** families, demonstrated antibacterial activity similar to the **HEA** family, except for polymers with a very low hydrophilic/hydrophobic ratio (**PEG-A/PEG-AA/HEA-I-1040**). Additionally, containing PEG as a neutral hydrophilic moiety exhibited significantly higher biocompatibility and selectivity compared to polymers with shorter, less hydrophilic, and less flexible chains (Table 5).

In a similar approach, Mortazavian *et al.*⁴⁵ adjusted the composition of cationic and hydrophobic monomers by introducing hydroxyethyl methacrylate (HEMA) monomer units to study the impact of each component on the antibacterial and hemolytic activities of the copolymer. The results showed that increasing the proportion of cationic groups in the copolymer beyond 30 mol% did not enhance its antibacterial efficacy against *E. coli*. They also proposed that the HEMA component (hydroxyl side chains) may have two active roles: (1) as a structural spacer to distribute the cationic and hydrophobic groups in monomer sequences and (2) as a modulator of copolymer chain insertion in bacterial and human cell membranes.

In summary, good neutral hydrophilic groups will maximize biocompatibility without negatively affecting the antibacterial efficacy of amphiphilic antimicrobial copolymers, which is of great significance for the design and synthesis of efficient and safe amphiphilic antimicrobial polymers.

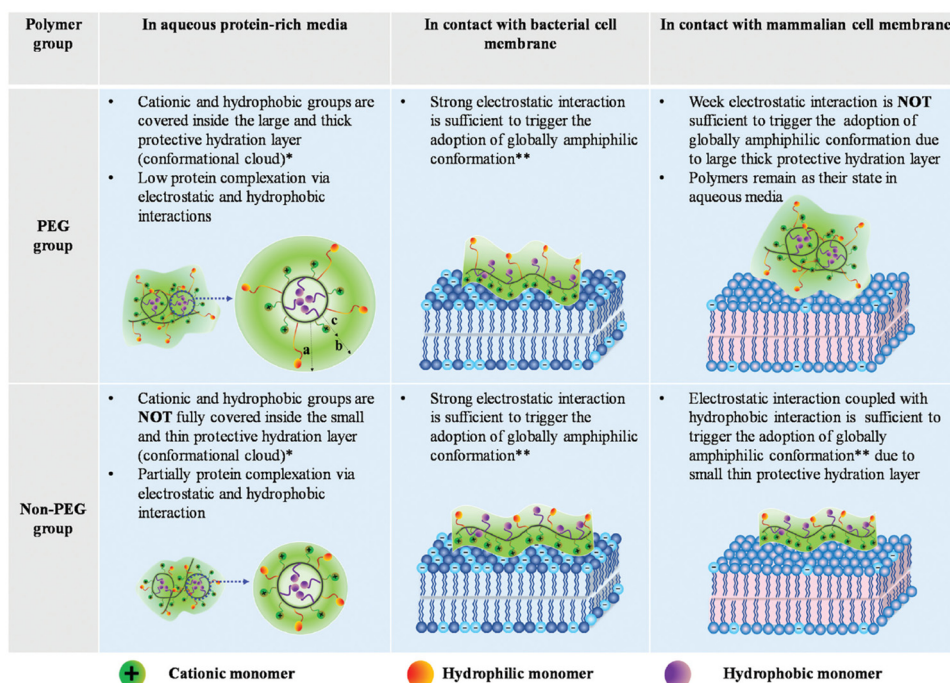


Fig. 9 Proposed adoption of PEG group polymer and non-PEG group polymer in protein-rich media and in contact with bacterial and mammalian cell membranes. Reproduced with permission from ref. 70 and 124, copyright 2010, Royal Society of Chemistry.



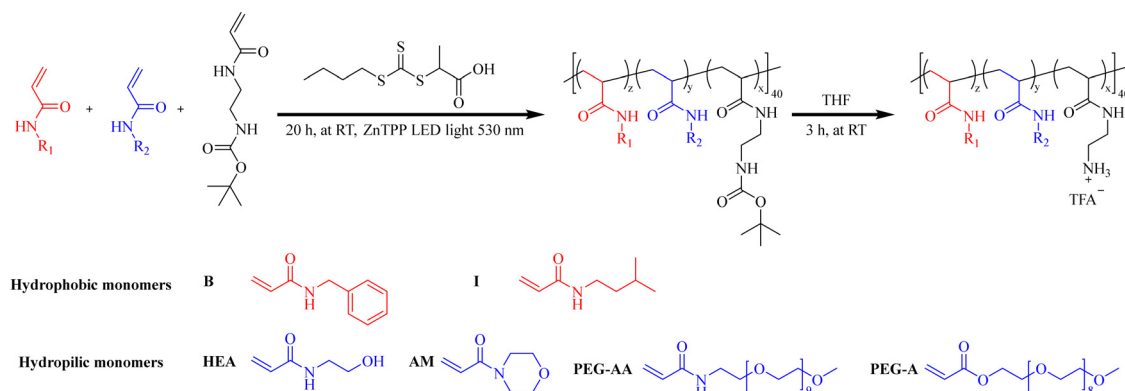


Fig. 10 Synthesis of amphiphilic antimicrobial copolymers using different hydrophobic and hydrophilic monomers. R1, R2 is hydrophobic or hydrophilic chain, respectively. Hydrophilic monomer: hydroxyethyl acrylamide (**HEA**), 4-acryloylmorpholine (**AM**), poly(ethylene glycol) methyl ether acrylate (**PEG-A**), and PEG acrylamide(**PEG-AA**), hydrophobic monomers: *N*-isopentylacrylamide (**I**) and *N*-benzylacrylamide (**B**). Redrawn from ref. 70.

2.1.3. Design of end group. As the polymerization degree and relative molecular mass ratio of the polymer decreases, the proportion of polymer end groups in the entire polymer increases. Therefore, the effect of end groups on polymer bioactivity is negligible for high molecular weight polymers but has a large effect for low molecular weight polymers or oligomers. For this type of amphiphilic antimicrobial copolymers, the relative molecular weight is mostly within 10 000, so the impact of end groups on their biological activity cannot be ignored.^{71,130,131} Additionally, advancements in RAFT polymerization have made it easier to control the synthesis of polymer chains with different end groups (Fig. 11).

Nadres *et al.*¹³⁰ used RAFT polymerization to produce a statistical methacrylate copolymer, followed by free radical-mediated conversion of the terminal groups successfully transformed the Z-end group of parent copolymer from dithiobenzoate

to a cyanoisobutyl or aminoethyl cyanopentanoate group without any significant changes to the polymer molecular weight. The results showed that the parent copolymer with dithiobenzoate end groups exhibited the strongest antibacterial and hemolytic activity, while the ability of copolymers to destroy membranes does not depend on the end group structure. In a more in-depth research, Griesser's group¹³¹ used three different RAFT chain transfer agents to systematically study the effects of changing the R- and Z-RAFT end groups on the antibacterial activity and cytotoxicity of poly(methacrylate) polymers (Fig. 12). The results indicate that the R-group predominantly influences the toxicity of the polymers. Substituting the anionic cyanovaleric acid R-group (**PA1**) with the neutral isobutyronitrile group (**PA3**) led to over a 20-fold increase in the hemolytic activity of the polymers. However, the Z-group has a greater impact on the antimicrobial activity of the polymers against MRSA and *C. albicans*, where polymers with

Table 5 Antimicrobial activities (MIC) and biocompatibility (HC₅₀ and IC₅₀) of 40-DP_n polymers with different hydrophobic and hydrophilic monomers and monomer feed ratios. Redrawn from ref. 70

Group	Family	Polymer	Monomer ratio (x:y:z)	MIC of polymer (μg mL ⁻¹)				HC ₅₀ (μg mL ⁻¹)	IC ₅₀ against MEF (μg mL ⁻¹)
				<i>P. aeruginosa</i> 01	<i>P. aeruginosa</i> 37	<i>E. coli</i> K12	<i>S. aureus</i> ATCC29213		
Non-PEG group	HEA	HEA-I-1040	50:10:40	16	—	16	>256	142 ± 6	—
		HEA-I-1535	50:15:35	16	32	16	>256	—	29 ± 4
		HEA-I-2030	50:20:30	16–32	32–64	32	>256	>2000	122 ± 19
		HEA-I-3020	50:30:20	>256	>256	>256	>256	>2000	—
		HEA-B-1040	50:10:40	16–32	—	32	256	961 ± 425	59 ± 37
		HEA-B-1535	50:15:35	32–64	—	64–128	>256	>2000	43 ± 3
	AM	AM-I-1535	50:15:35	8–16	16–32	16	>256	1825	56 ± 13
		AM-I-2030	50:20:30	16	32	16	>256	1505 ± 29	56 ± 15
		AM-I-3020	50:30:20	64	64	64–128	>256	>2000	326 ± 101
		AM-B-1040	50:10:40	32	32–64	32–64	—	878 ± 81	39 ± 25
		AM-B-1535	50:15:32	32	32–64	32–64	—	855 ± 212	36 ± 7
		AM-B-1535	50:15:32	32	32–64	32–64	—	855 ± 212	36 ± 7
PEG group	PEG-AA	PEG-AA-I-1040	50:10:40	8	16–32	8–16	>256	390 ± 87	43 ± 10
		PEG-AA-I-1535	50:15:35	16	32	8–16	>256	>2000	166 ± 17
		PEG-AA-I-2030	50:20:30	16–32	32–64	32	>256	>2000	~512
	PEG-A	PEG-A-I-1040	50:10:40	8	16–32	8–16	>256	500 ± 343	39 ± 2
		PEG-A-I-1535	50:15:35	16	32–64	8–16	>256	>2000	130 ± 30
		PEG-A-I-2030	50:20:30	16–32	64–128	32–64	>256	>2000	~512
		PEG-A-I-3020	50:30:20	>256	>256	>256	>256	>2000	>512
		PEG-A-B-1040	50:10:40	16–32	32–64	32–64	—	1563 ± 135	56 ± 7
		PEG-A-B-1535	50:15:35	32–64	64	64–128	—	>2000	191 ± 30

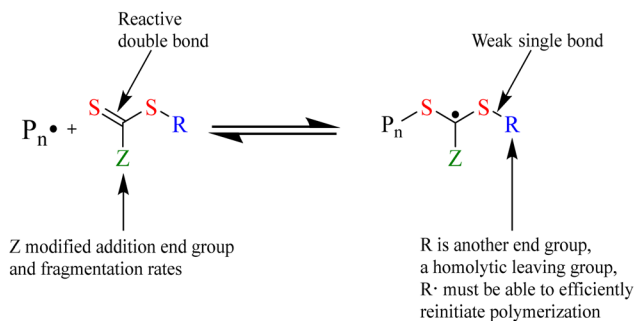


Fig. 11 Scheme of RAFT polymerization's main mechanism step.⁴⁶

a long, lipophilic dodecylsulfanyl Z-group (**PA1**) were more effective than those with either an ethylsulfanyl or no Z group.

In conclusion, although the end group is not the main component of amphiphilic antimicrobial copolymers, modulating the end group should not be ignored to achieve optimal amphiphilic antimicrobial copolymers.

2.2. Global polymer design

2.2.1. Molecular weight. The association between molecular weight and antibacterial effectiveness seems to be more

pronounced in Gram-positive bacteria compared to Gram-negative ones.^{51,72,132} Specifically, Gram-positive bacteria feature a densely interconnected peptidoglycan layer that serves as a barrier, shielding the inner membrane from larger membrane-active molecules. This phenomenon, termed the “sieving effect,” was initially recognized by Lienkamp *et al.*¹³³ and subsequently corroborated by numerous independent investigations. For instance, Pachla and co-workers¹³⁴ studied the effect of molecular weight on the antibacterial, cytotoxic and hemolytic activities of linear polytrimethyleneimine (L-PTMI). The results (Table 6) showed that within the test range (0.8–18 kDa), the antibacterial activity of the polymer increased with increase in polymer molecular weight. Above a certain molecular weight, activity increases significantly, manifested by lower MIC, HC₅₀, and IC₅₀ values. Notably, the threshold for MIC values appears to be at a lower M_n or M_w compared to HC₅₀ and IC₅₀ values. Furthermore, Qiao *et al.*¹³⁵ reported the antibacterial activity of a series of random polycarbonate polymers synthesized *via* metal-free organocatalytic ring-opening polymerization. Their study revealed a positive correlation between antibacterial activity against *Staphylococcus aureus* and molecular weight. Polymers with molecular weights of 6 kDa, 10 kDa, and 16 kDa (each with a charge density of approximately 50%)

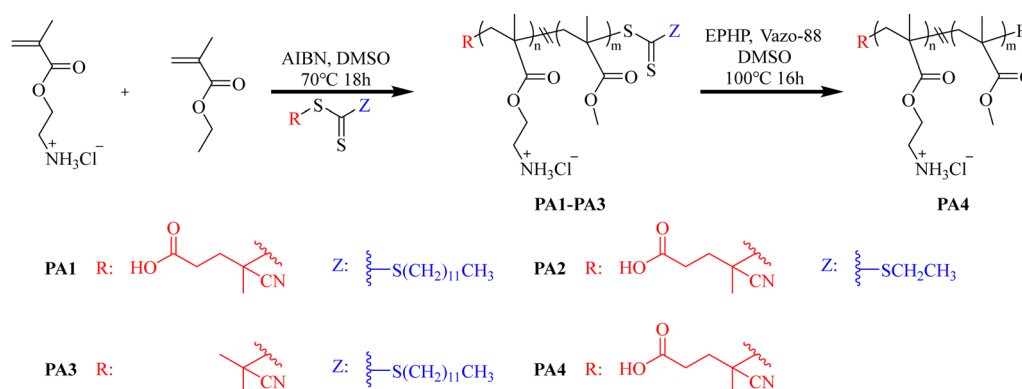


Fig. 12 Synthesis of amphiphilic antimicrobial copolymers (**PA1–PA4**) by using RAFT agents to control the variation in the Z and R groups at the polymer ends. Redrawn from ref. 131.

Table 6 Biological activity and molecular weight of the studied linear polytrimethyleneimine (L-PTMI). Redrawn from ref. 134

Polymer	M_n /kDa	M_w /kDa	PDI	MIC/ $\mu\text{g mL}^{-1}$			HC ₅₀ / $\mu\text{g mL}^{-1}$	IC ₅₀ / $\mu\text{g mL}^{-1}$
				<i>E. coli</i>	<i>S. aureus</i>	<i>C. albicans</i>		
L-PTMI_0.8k	0.80	0.91	1.14	256	128	32	> 2000 (2%)	2220
L-PTMI_1.2k	1.2	1.4	1.22	16	16	16	> 2000 (4%)	2570
L-PTMI_3k	2.8	3.7	1.33	2	4	16	> 2000 (3%)	25.2
L-PTMI_7k	6.7	9.6	1.44	2	1	16	125	4.20
L-PTMI_12k	12.1	18.2	1.51	2	1	32	15.6	2.48
L-PTMI_18k	18.0	27.1	1.51	2	1	32	9.40	2.71
L-PEI_4k	4.0	5.3	1.30	16	16	32	> 2000 (39%)	660
Me-L-PTMI_7k	3.6	5.4	1.53	16	16	16	> 2000 (2%)	240
MePTMI-co-Me ₂ PTMI _{10%}	4.2	6.5	1.54	16	16	16	> 2000 (3%)	420
MePTMI-co-Me ₂ PTMI _{20%}	3.7	5.5	1.49	16	16	16	> 2000 (3%)	930
Me-L-PEI_4k	3.9	4.8	1.24	32	128	64	> 2000 (2%)	600
MePEI-co-Me ₂ PEI _{10%}	3.6	4.6	1.27	32	64	32	> 2000 (3%)	1370
MePEI-co-Me ₂ PEI _{20%}	3.6	4.7	1.31	16	64	16	> 2000 (2%)	2000

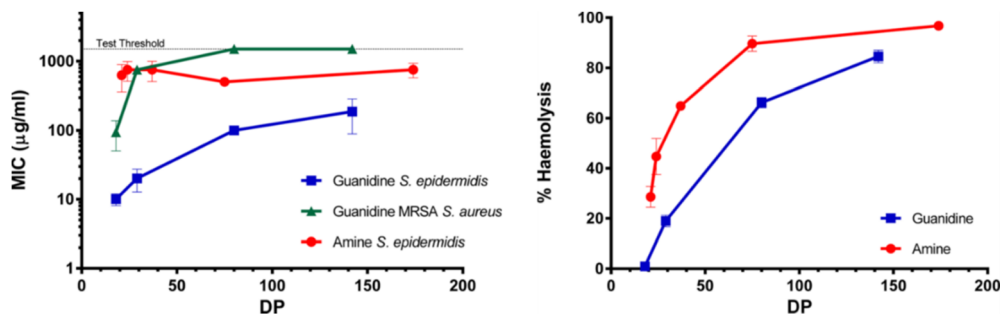


Fig. 13 (a) Antimicrobial activities (MIC) and (b) hemotoxicity of guanidine and amine polymers as a function of polymer chain length (DP). Reproduced with permission from ref. 51, copyright 2013, American Chemical Society.

exhibited MIC values of 63 mg L^{-1} , 125 mg L^{-1} , and 500 mg L^{-1} , respectively. The authors attributed this to the “sieving effect,” wherein higher molecular weight polymers are captured by the dense peptidoglycan layer on the outer surface of *S. aureus*, limiting their effectiveness. Locock *et al.*⁵¹ conducted a comparative study on the antibacterial and hemolytic activities of cationic polymethacrylates containing either amine or guanidine groups and hydrophobic side chains. Their findings revealed that an eightfold increase in polymer molecular weight resulted in minimal changes in the MIC values of amine-based polymers, whereas a similar increase in guanidine-based copolymers caused a nearly 20-fold rise in MIC values (Fig. 13(a)). The authors hypothesized that this discrepancy arises due to potential differences in the mechanisms of action between amine and guanidine copolymers. Furthermore, for both guanidine and amine copolymers, increasing the molecular weight led to heightened hemolytic activity (Fig. 13(b)). Notably, the shortest guanidine polymer chains exhibited negligible hemolytic activity at their MIC values, while the longest chains showed significant hemolytic activity under the same conditions. Nevertheless, guanidine copolymers consistently demonstrated lower hemolytic activity at their MIC values compared to their amine counterparts, indicating better biocompatibility for guanidine-based systems.

Therefore, amphiphilic antimicrobial copolymers with different effects on Gram-positive and Gram-negative bacteria can be designed and synthesized by adjusting the molecular weight to meet the needs of different users.

2.2.2. Topology of amphiphilic antimicrobial copolymers.

The topology of polymers refers to the spatial arrangement of monomers along their chains. Recent advancements in polymer chemistry have led to the development of numerous antimicrobials with various topological structures.^{46,136,137} These range from simple structures like linear and cyclic polymers to advanced structures such as hyperbranched, dendritic star-shaped, or brush-like polymers. These diverse structures significantly influence the physical, chemical, and biological properties of polymers.

2.2.2.1. Linear polymers. Linear polymers have been extensively studied due to their fundamental topological structure. An array of amphiphilic antimicrobial copolymers can be produced through diverse active polymerization techniques.

The progress in controlled polymerization methodologies allows for precise adjustment of parameters like molecular weight, component composition ratios, and topological architectures, encompassing homopolymers, random, alternating, and block copolymers.^{21,138,139}

2.2.2.1.1. Random polymers. Random polymers represent the simplest topology among copolymers. It is notable that numerous research reports have proven that random amphiphilic copolymers have efficient antibacterial effects.^{51,53,68,72,117,128,140–142} Due to their straightforward synthesis, devoid of any requirement for precise sequence control, the simplicity of the technology combined with favorable outcomes has made random copolymers an important part of the amphiphilic antimicrobial copolymers field.

However, the spatial distribution of components is bound to affect the secondary structure of the polymer, thereby affecting the overall biological efficacy by affecting its physicochemical properties and the conformation of the biological interface. Inspired by natural HDPs, researchers are dedicated to creating precisely defined sequence amphiphilic antimicrobial copolymers, manipulating the concentration of local structural domains within polymer chains, and mimicking the precise monomeric sequences and secondary sequences found in natural HDPs. For example, alternating or block copolymers are designed to enhance their biological activity and improve their selectivity between bacteria and mammalian cells.^{143–145}

2.2.2.1.2. Block polymers. Block polymers are the most widely reported type of well-defined sequence.^{80,146–149} They are usually synthesized by living polymerization technologies, such as ATRP and RAFT polymerization.^{46,150}

Many research groups^{142,151–153} have reported the preparation of random and diblock copolymers containing the same structure and composition and compared their biological activities. The results both showed that random and diblock copolymer series displayed similar antibacterial activity. But the random copolymer showed very high hemolytic activity and very low cell selectivity, while the diblock copolymer showed significantly lower hemolytic activity and significantly higher cell selectivity. A widely accepted hypothesis is that this is related to the difference in single chain conformation between block copolymers and random copolymers (Fig. 14). Block copolymers can form intramolecular aggregates surrounding a hydrophobic core of cationic segments in water. The



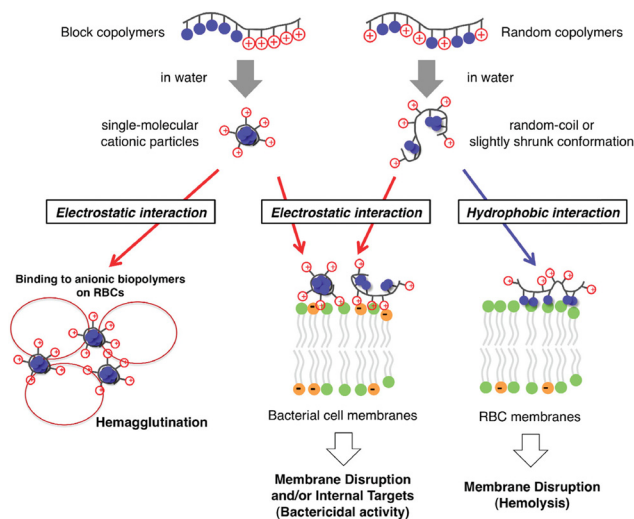


Fig. 14 Schematic illustration of the antibacterial and hemolytic activities of random and block amphoteric copolymers. Reproduced with permission from ref. 151, copyright 2011, American Chemical Society.

cationic nature of these intramolecular aggregates increases their electrostatic binding to the anionic LPS on the bacterial surface and may displace the divalent cations that stabilize the outer membrane structure. Replacement of these divalent cations disrupts the membrane structure and increases the permeability of the outer membrane, thus facilitating the entry of polymers into the cell surface. As for the hemolytic activity, the hydrophobic segment was shielded by the cationic surface, which reduced the hydrophobic binding of the polymer to the cell membrane of RBCs and, therefore, did not cause significant hemolytic activity.¹⁵³ On the other hand, the random copolymers, will adopt a random coil or slightly shrunk conformation because the hydrophobic groups randomly distributed along the polymer chain may not form a hydrophobic

core if the hydrophobic content is low enough. These hydrophobic groups interact with the red blood cell membrane and cause cell lysis. In addition, the cationic functional groups of random copolymers can also enhance the binding of the polymer to bacterial cell membranes and cause membrane rupture.^{126,154}

2.2.2.2. Cyclic polymers. Another possible structure for amphiphilic antimicrobial copolymer is a cyclic one. Cyclic polymers are linked end to end to form a ring structure and therefore have no end groups.¹⁵⁵ Despite being inspired by many natural cyclic structures of amphiphilic antimicrobial copolymers,¹⁵⁶ the utilization of cyclic amphiphilic polymers for antimicrobial applications remains relatively underexplored compared to other topological types. This may be attributed to the considerable difficulty in obtaining highly purified cyclic polymers.

For example, Xu and co-workers¹⁵⁷ prepared cyclized cationic copolymers through ATRP (Fig. 15). These copolymers are more readily attracted to negatively charged bacterial cell membranes and cause membrane damage, exhibiting better antimicrobial performance compared to their linear counterparts. Additionally, the cytotoxicity of the cyclic cationic copolymers is slightly lower than that of the linear copolymers.

Aquib *et al.*¹⁵⁸ synthesized a functional RAFT agent containing benzaldehyde groups (Fig. 16(A)), which was subsequently utilized to prepare amphiphilic cationic statistical linear terpolymers *via* RAFT polymerization. These linear terpolymers were then cyclized into cyclic terpolymers through a Hetero-Diels-Alder click reaction (Fig. 16(B)). Compared to their linear counterparts, the cyclic terpolymers exhibited superior hemocompatibility and enhanced biocompatibility with mammalian macrophages, demonstrating improved safety profiles.

This indicates that cyclic polymers hold significant research value in the field of amphiphilic antimicrobial copolymers. It is

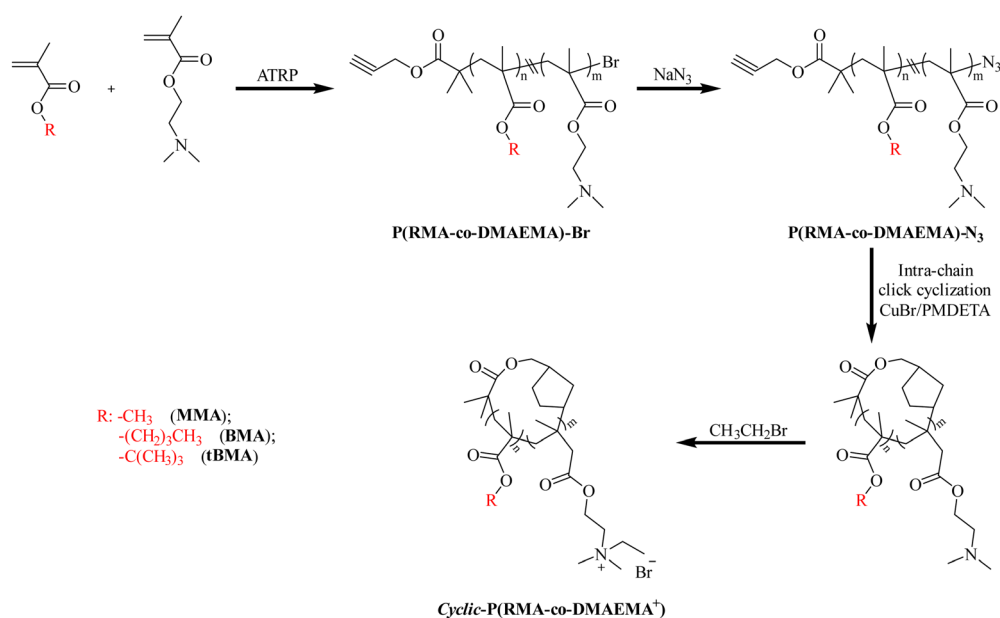


Fig. 15 Synthesis of cyclic cationic P(RMA-co-DMAEMA⁺) copolymers with different hydrophobic groups. Redrawn from ref. 157.

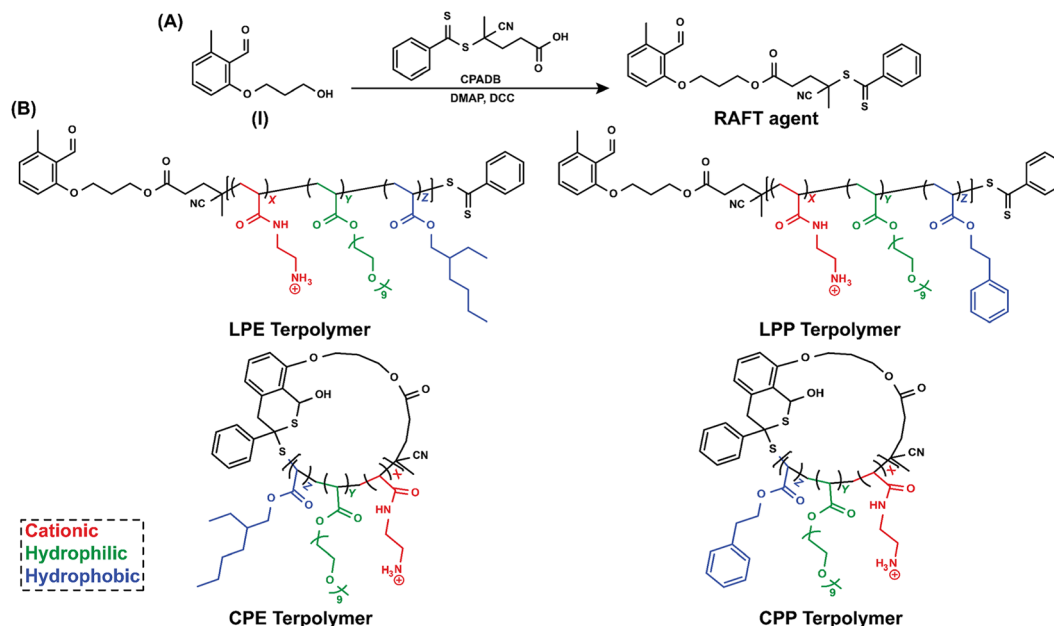


Fig. 16 (A) Synthesis scheme of RAFT agent. (B) Chemical structures of amphiphilic LPE, CPE, LPP, and CPP (X = percentage of cationic groups, Y = percentage of hydrophilic groups, and Z = percentage of hydrophobic groups). Redrawn from ref. 158.

anticipated that with innovations in the synthesis methods of cyclic polymers,^{159,160} more research on cyclic amphiphilic polymers will emerge in the future.

2.2.2.3. Advanced architecture design. Inspired by the precise and complex protein structure, many researchers tried to develop amphiphilic antimicrobial copolymers with higher structures. Along with a deeper understanding of amphiphilic antimicrobial copolymers, advances in polymer chemistry have made it possible to artificially synthesize a variety of advanced polymers with complex structures, such as hyperbranched, dendritic star or brush polymers.

For example, Santos *et al.* reported¹⁶¹ the synthesis of linear, 4-arm star and 6-arm star copolymers *via* supplemented activator and reducing agent atom transfer radical polymerization (SARA-ATRP) (Fig. 17). The results showed that 6-arm and 4-arm star polymers with similar molecular weights exhibited stronger antibacterial activity than linear polymers.

Aquib *et al.*¹⁶² synthesized linear polymers (LPs), hyperbranched polymers (HPs) and star polymers (SPs) *via* RAFT polymerization (Fig. 18). The results showed that the blood compatibility of HPs with the same ratio of hydrophobic to hydrophilic groups was significantly improved by 2 to 4 times compared with that of LPs; SPs with a low hydrophobic ratio are not toxic to red blood cells and retain similar potent antibacterial activity to linear polymers and hyperbranched polymers. Overall, SPs exhibit superior biological activity compared with HPs and LPs (SPs > HPs > LPs).

Laroque *et al.*¹⁶³ successfully utilized an arm-first synthesis strategy to create a library of amphiphilic cationic core-cross-linked star (CCS) copolymers through RAFT polymerization (Fig. 19). These polymers featured both diblock and homopolymer arms. The results indicated that transitioning from diblock to miktoarm CCS structures enhanced their antibacterial activity. Furthermore, while none of the synthesized polymers exhibited hemolytic activity in sheep red blood cells, there

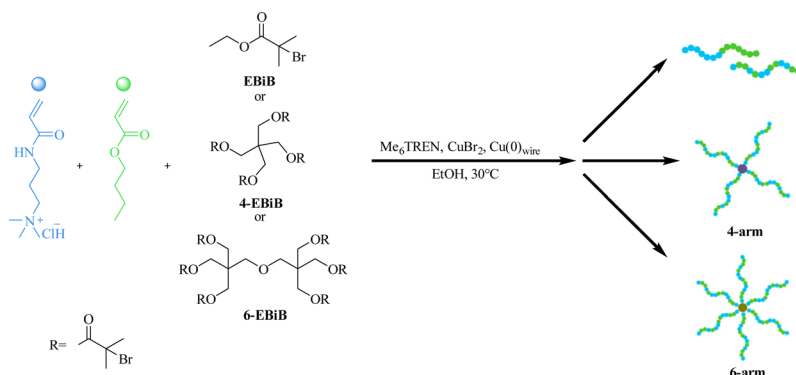


Fig. 17 Synthesis of the amphiphilic linear and linear, 4-arm star and 6-arm star shaped copolymers by SARA-ATRP. Redrawn from ref. 161.

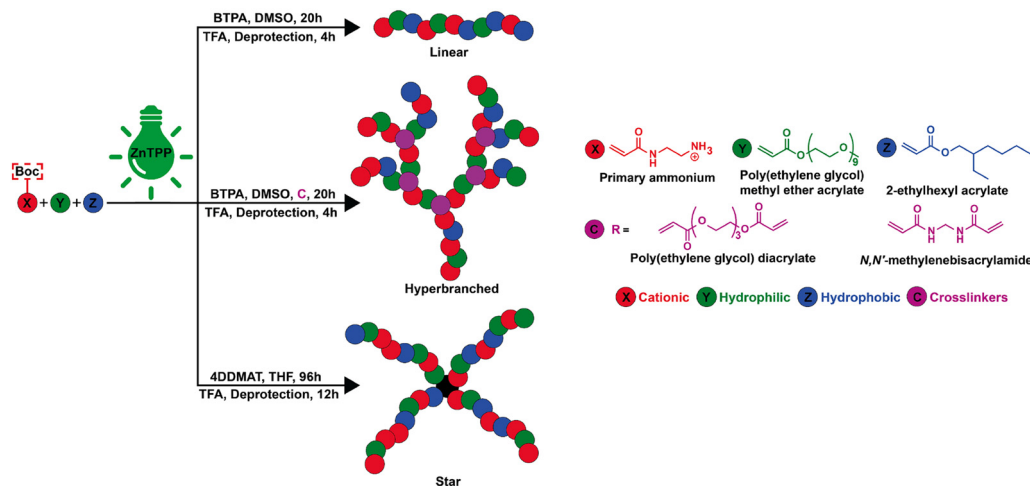


Fig. 18 Synthesis of LPs, HPs, and 4-arm SPs prepared by RAFT polymerization. Redrawn from ref. 162.

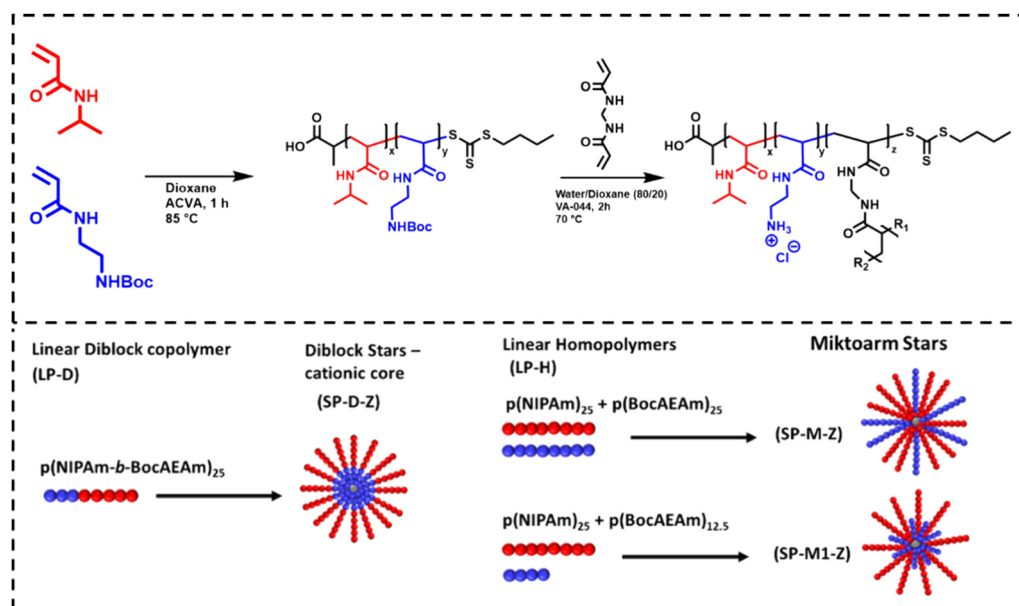


Fig. 19 Planned synthesis and nomenclature of the "arm-first" core-cross-linked star (CCS) copolymer library. Linear polymers are labeled LP-X (LP = linear polymer, four different polymers numbered X = 1–4) and star polymers SP-D/M-Z (SP = star polymer, D = diblockarm, M = miktoarm, Z = DP of cross-linker). Redrawn from ref. 163.

was a notable increase in the tendency for red blood cell aggregation with greater exposure of cationic units. Importantly, increasing the number of arms did not improve antibacterial activity compared to 4-arm polymers. The authors attributed this to the shielding effect of cationic units located in the polymer core, which hindered membrane interactions, although the miktoarm structures showed slight improvements in this regard.

Islam *et al.*¹⁰³ synthesized dendritic amphiphilic antimicrobial copolymers through ROMP, that showed strong cell selectivity, with high antibacterial activity against Gram-positive bacteria (*S. aureus*) ($\text{MIC} = 32 \mu\text{g mL}^{-1}$) and non-toxicity to RBCs ($\text{HC}_{50} > 1000 \mu\text{g mL}^{-1}$). However, all polymers were

inactive against Gram-negative bacteria (*E. coli*) ($\text{MIC} > 256 \mu\text{g mL}^{-1}$). They hypothesize that dendritic copolymers, each containing three positive charges per dendritic unit, along with hexyl pyridinium functional groups, adhere to the lipopolysaccharide and outer membrane layers, resulting in diminished activity against *E. coli*.

Up to now, many studies have reported that amphiphilic antimicrobial copolymers with complex structures, including hyperbranched,^{162,164} dendritic star^{161,165} or brush polymers^{166,167} exhibit strong antibacterial activity and better biocompatibility than their linear counterparts. One potential explanation may be attributed to their intrinsic multivalent compact structures, where numerous end groups provide high local concentrations of active

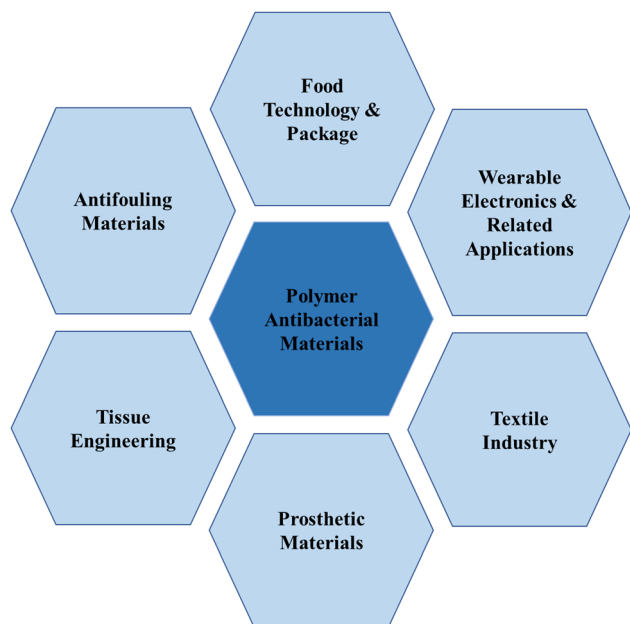


Fig. 20 Some applications of polymer antibacterial materials.

functional groups. These active moieties may synergistically enhance the initial adsorption and interaction between complex macromolecules and bacterial cells, subsequently leading to increased membrane disruption.

3. Antimicrobial applications of amphiphilic copolymers

In recent years, amphiphilic antimicrobial polymers have attracted much attention due to their low cost, simple preparation, effective antibacterial effect, easy modification, good biocompatibility, safety and non-toxicity. These properties makes amphiphilic antimicrobial copolymers highly desirable for a wide range of applications, particularly in the biomedical field, textiles, agriculture, food packaging, building materials, water treatment industry, public facilities, and medical facilities and equipment^{168,169} (Fig. 20). The next part of this review will discuss the application prospects of this type of amphiphilic antimicrobial polymers in different aspects, including coatings, hydrogels, vesicles, antibiotic substitutes, and more.¹⁷⁰

3.1. Coatings

Material surfaces with antibacterial properties can prevent bacteria from adhering on and colonizing the surface of objects to form biofilms. The simplest way to give antibacterial functions to the surface of materials is to coat them. An ideal antimicrobial coating should meet the following main requirements: (a) be bactericidal against a broad spectrum of pathogenic microorganisms; (b) be active for a long time (preferably permanently); (c) be stable (should not decompose into toxic products); (d) be environmentally friendly; (e) be easy and cheap to synthesize; (f) be insoluble in water in many

applications.^{171–173} Some type of synthetic amphiphilic antibacterial polymers meet these requirements.

For example, Ma *et al.*¹⁷⁴ designed a multifunctional environmentally adaptable polymer coating based on poly methacryloxyethyl dimethyl butyl ammonium bromide–polydimethylsiloxane (pMDBAB–PDMS) modified PET film (Fig. 21(a)). The soft polysiloxane chains and hydrophilic quaternary ammonium molecules enable the coating to have optimal anti-fog, antibacterial, antistatic, self-cleaning and self-lubricating properties in both water and air. At the same time, the cationic quaternary ammonium groups of the pMDBAB unit are covalently bonded in the coating network, making the coating highly antibacterial against Gram-positive *S. aureus* and Gram-negative *E. coli*. In addition, since the PDMS chains preferentially assemble in the air, the coating also exhibits good lubrication and oil adhesion resistance under dry air conditions.

Bai *et al.*¹⁷⁵ used synthetic polyhedral oligomeric silsesquioxane–poly(quaternary ammonium compound-co-2-aminoethyl methacrylate hydrochloride) (POSS–P(QAC-co-AEMA)) and poly(*N*-hydroxyethylacrylamide-co-glycidyl methacrylate) (P(HEAA-co-GMA)) to coat endoscopic lenses used in surgery (Fig. 21(b)). Due to the high water absorption capacity of HEAA and QAC, the coating has excellent anti-fogging properties under both *in vitro* and *in vivo* (mouse) fogging conditions. And by simply adjusting the mixing ratio of POSS–P(QAC-co-AEMA) and P(HEAA-co-GMA), the coating can achieve comprehensive bacteria-killing and bacteria-repelling properties, as well as a lower hemolysis rate.

Wang *et al.*⁸⁰ synthesized a series of triblock copolymers with bactericidal quaternary ammonium groups at both ends of the polymer chain and bio-responsive PCL segments in the polymer chain (Fig. 21(c)). This triblock copolymer readily self-assembles to form RM with hydrophobic PCL segments in the shell and bactericidal quaternary ammonium groups in the core. Benefiting from the good biocompatibility of PCL, natural RM has little cytotoxicity. They demonstrated that RM can be easily impregnated into commercial gelatin sponge to fabricate an autonomously responsive antimicrobial coating (RM coated gelatin sponge) for the engineering design of self-sterilizing implantable medical dressings. *In vivo* experimental results in mice showed that when lipase emerges from released at the site of infection, PCL fragments are hydrolyzed, causing quaternary biocidal agents to be released in response, killing bacteria in the vicinity of the implant. In particular, *S. aureus* and *B. subtilis* were susceptible to quaternary biocidal agents released from RM-coated gelatin sponge.

Chen *et al.*¹⁷⁶ employed a synthesized random terpolymer consisting of glycidyl methacrylate-derived quaternary ammonium cations (G-MPA-N⁺) as the antimicrobial component, zwitterionic 2-methacryloyloxyethyl phosphorylcholine (MPC) as the fouling-resistant component, and dopamine methacrylamide (DMA) as the anchoring moiety to modify the activated surface of PDMS elastomer, thereby constructing a dual-functional antibacterial coating (Fig. 21(d)). The resultant dual-functional antibacterial coating exhibits biocompatibility, effectively inhibiting the formation of pathogenic biofilms. Furthermore, it demonstrates



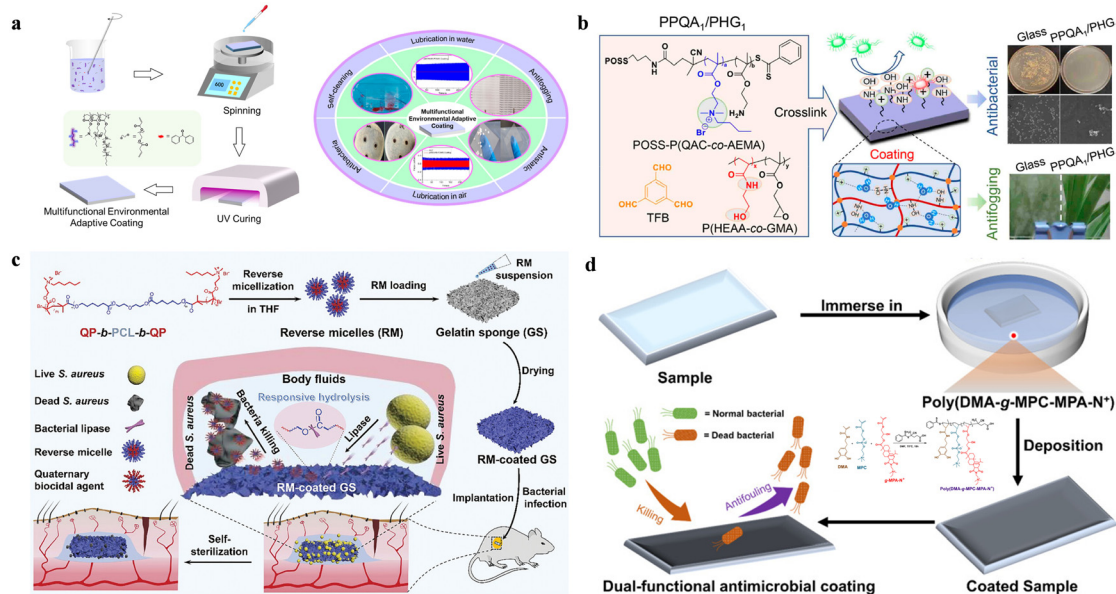


Fig. 21 (a) Preparation process of amphiphilic multifunctional environmental adaptive coating; reproduced with permission from ref. 174, copyright 2022, American Chemical Society. (b) Synthesis of POSS-P(QAC-co-AEMA) and P(HEAA-co-GMA) copolymers for antibacterial and antifogging coatings; reproduced with permission from ref. 175, copyright 2020, American Chemical Society. (c) Schematic illustration for the fabrication of RM-coated GS and the mechanism of bioswitchable self-sterilization in bacterial infection site; reproduced with permission from ref. 80, copyright 2021, Wiley. (d) Preparation process of poly(DMA-G-MPC-MPA-N⁺) and dual-functional antimicrobial coating; reproduced with permission from ref. 176, copyright 2022, Elsevier.

minimal cytotoxicity towards mammalian cells concurrently. Of note, in an *in vivo* implantation model, the antibacterial coating can eradicate pathogenic biofilms within implants, preventing host tissue damage and inflammation.

These studies indicate that using this amphiphilic antimicrobial polymer as a coating has great application prospects, especially in the field of surgical medical devices, and it is believed that this research field will be further developed.

3.2. Hydrogels

Hydrogels are three-dimensional polymer networks that contain large amounts of water through hydrophilic groups or domains that hydrate in an aqueous environment.¹⁷⁷ Chemical or physical cross-linking can render the network of homopolymers or copolymers resistant to dissolution in the aqueous phase. In addition to conventional properties such as high-water content, tunable mechanical properties, and lack of cytotoxic products after degradation, hydrogels also exhibit excellent biocompatibility and bioactivity.^{178,179} Due to the unique properties of antimicrobial hydrogels, they are widely used in biomedicine, tissue engineering, and smart textiles, including applications in drug delivery, regenerative medicine, wound dressings, and *in vitro* diagnostics. Antimicrobial hydrogels based on amphiphilic antimicrobial copolymers have also received significant attention because they combine the advantages of hydrogels and additives to exploit multiple properties on a single platform.^{180,181} For instance, the porous structure of hydrogels allows water/blood to flow in quickly, giving the material instant blood absorption capabilities and can also act as a barrier to prevent blood loss by forming a hemostatic plug.^{182,183} Therefore, the application of amphiphilic antimicrobial

copolymers loaded hydrogels in wound healing dressings is a promising strategy.^{184–186} Particularly interesting are also injectable hydrogels, that can fill up wounds and cross-link *in situ*, combining antimicrobial activity with hemostatic capacity.¹⁷³

To address specific application scenarios, as illustrated below (Fig. 22(a)), a biodegradable amphiphilic block copolymer was utilized to design a temperature-responsive antimicrobial hydrogel.¹⁸⁴ At room temperature, these systems exist as soluble micelles in aqueous solution. However, at physiological temperature (approximately 37 °C), they transform into gel-like materials with distinctive supramolecular fiber/ribbon-like structures and exhibit shear-thinning behavior. This hydrogel demonstrates broad-spectrum antimicrobial activity and excellent skin biocompatibility.

Zhao *et al.*¹⁸⁵ utilized the spontaneous radical polymerization of [2-(methacryloyloxy)ethyl]dimethyl-(3-sulfopropyl) ammonium hydroxide (SBMA) to prepare an arginine-modified chitosan-oligosaccharide (COS-Arg) doped hydrogel system *via* a simple one-pot method (Fig. 22(b)). SBMA was covalently cross-linked with DMA and methacrylateoethyl trimethyl ammonium chloride (DMC) to form the hydrogel framework. The hydrogel exhibited excellent antibacterial activity and superior cytocompatibility. Additionally, it maintained a moist microenvironment at the wound site, promoted the production of vascular endothelial growth factor and hydroxyproline, reduced the formation of tumor necrosis factor- α , and effectively accelerated wound healing.

Additionally, Wang *et al.*¹⁸⁷ encapsulated nanoengineered peptide-grafted hyperbranched polymers (NPGHP) in an anionic polyelectrolyte and, for the first time, constructed

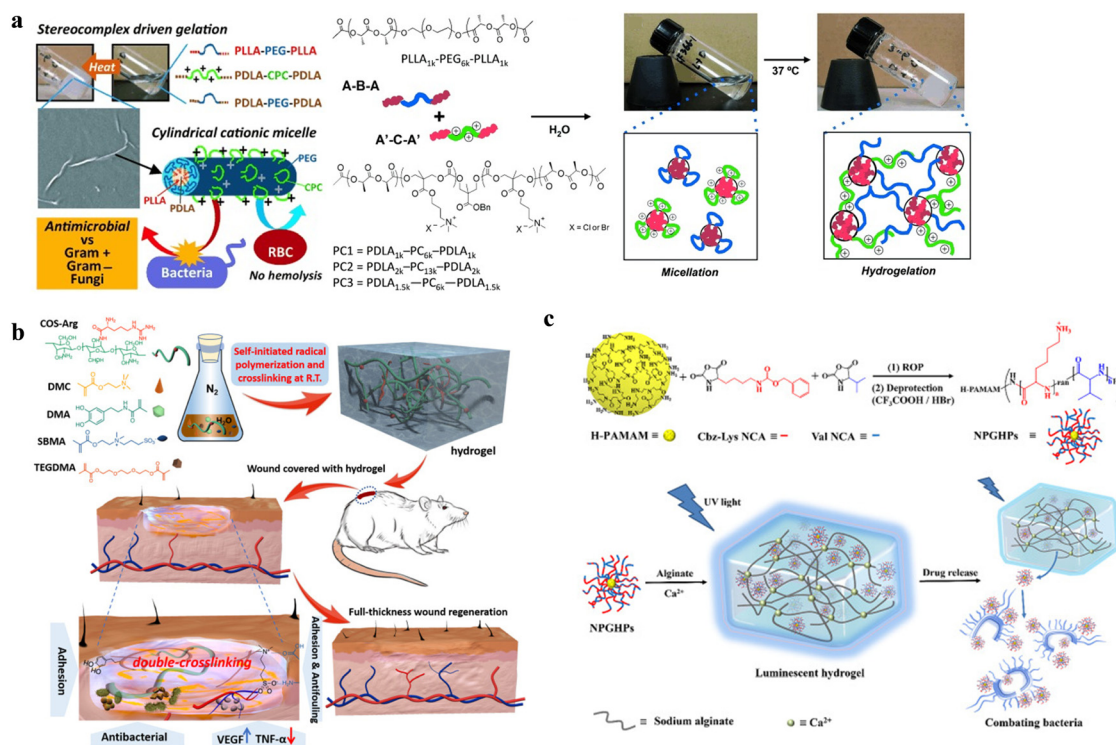


Fig. 22 (a) Preparation of temperature-sensitive mixed micellar solutions and gels containing PLLA-PEG-PLLA and PDLA-CPC-PDLA; reproduced with permission from ref. 184, copyright 2013, Wiley. (b) Schematic diagram of one-pot self-initiated radical polymerization to prepare highly adhesive and double-crosslinked antibacterial and antifouling gel dressing and its application in wound healing; reproduced with permission from ref. 185, copyright 2022, Elsevier. (c) Schematic diagram of the synthesis route of NPGHPs and their application in visual monitoring and antibacterial NPGHPs/SA gel; reproduced with permission from ref. 187, copyright 2022, Elsevier.

aggregation-induced emission (AIE) active fluorescent polymeric hydrogels (NPGHP/sodium alginate (SA) gels) using non-conjugated luminescent polymers (NCLPs) with AIE characteristics (Fig. 22(c)). The NPGHP/SA gel system, with good biocompatibility, exhibited significant luminescence changes during the release process, thereby enabling the visualization of dynamic drug release behavior. This system provides an advanced biomedical material with broad-spectrum antibacterial properties and offers a facile method for investigating drug release in hydrogel systems.

Distinct from common polymer hydrogels, Brahmachari *et al.*¹⁸⁸ designed a novel amphiphilic hydrogelator utilizing physical cross-linking, such as hydrogen bonding between compounds. This design involved connecting a quaternary pyridinium unit to a hydrophobic long alkyl chain *via* an amide bond, resulting in a compound with a very simple synthesis. Efficient hydrogel formation was achieved through the interdigitated bilayer packing of the amphiphilic molecules, leading to the development of three-dimensional supramolecular aggregates. These hydrogelators exhibited high antibacterial activity against both Gram-positive and Gram-negative bacteria ($\text{MIC} < 0.4 \mu\text{g mL}^{-1}$) while demonstrating very low cytotoxicity ($\text{IC}_{50} > 100 \mu\text{g mL}^{-1}$).

There are also many studies that have investigated the incorporation of antimicrobial agents into hydrogels to achieve antimicrobial effects.^{189–192} However, wound dressings

containing antimicrobial agents may lose their long-term bactericidal efficacy due to the leaching of these agents. Furthermore, certain nano-inorganic metal-based antimicrobial agents exhibit cytotoxicity, posing potential risks to biological systems.^{193–195} Additionally, traditional hydrogel systems often utilize initiators such as ammonium persulfate (APS) or 2-hydroxy-4'-(2-hydroxyethoxy)-2-methylpropiophenone (HHMP), which can leach from the hydrogel matrix. This leaching may irritate and corrode mucosal tissues and, with prolonged skin exposure, lead to allergic dermatitis.¹⁹⁶ Consequently, the development of inherently antimicrobial hydrogels that do not rely on toxic initiators has emerged as a promising research direction. Such hydrogels offer the dual advantages of sustained antibacterial efficacy and excellent biocompatibility, addressing the limitations of traditional hydrogel systems.

3.3. Vesicles

Typically, polymer vesicles are nanoscale hollow spheres composed of three parts: an inner cavity, a hydrophobic membrane, and a hydrophilic corona, making them more complex than polymer micelles with simple core-shell nanostructures.¹⁹⁷ Amphiphilic antimicrobial copolymers with cationic groups and hydrophobic chains (often as nanoparticles) have demonstrated excellent antibacterial performance against both Gram-positive and Gram-negative bacteria.^{198,199} However, strong positive charges can lead to



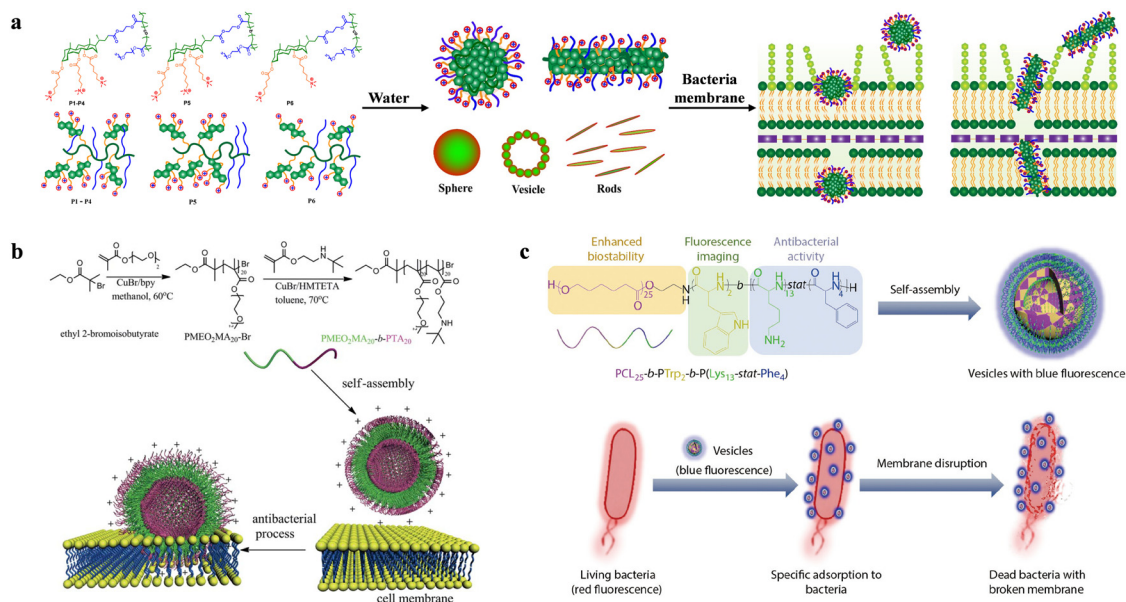


Fig. 23 (a) Schematic diagram of the antibacterial mechanism of the designed and synthesized cholic acid based gradient copolymers (**P1–P4**), diblock copolymer (**P5**), and deoxycholic acid based copolymer (**P6**) on the cell membrane of Gram-negative bacteria; reproduced with permission from ref. 198, copyright 2022, American Chemical Society. (b) Schematic diagram of antimicrobial vesicles formed by directly dissolving diblock copolymers in water; reproduced with permission from ref. 147, copyright 2013, Royal Society of Chemistry. (c) Synthesis of peptide-based vesicles with antibacterial activity and intrinsic blue fluorescence, and schematic diagram of real-time visualization of the antibacterial process; reproduced with permission from ref. 204, copyright 2021, Springer Nature.

severe cytotoxicity, such as hemolysis, which is a common side effect of cationic polymers. Therefore, forming nanostructured vesicles with amphiphilic antimicrobial copolymers that possess moderate positive charges may be a comprehensive strategy to balance antibacterial activity and cytotoxicity.^{91,147,151,200–203}

For instance, Rahman *et al.*¹⁹⁸ selected bile acid and neutral PEG components to fabricate amphiphilic antimicrobial copolymers capable of forming nanoscale particles. These amphiphilic antimicrobial copolymers form spheres, rods, and vesicles (Fig. 23(a)). The nanoscale structures of these bile acid-based macromolecules preferentially interact with bacterial membranes, exhibiting broad-spectrum antibacterial activity against both Gram-positive and Gram-negative bacteria. The cholic acid-based copolymers with three QAC charges (**P1–P5**) demonstrated better efficacy compared to the deoxycholic acid-based copolymers with dual QAC charges (**P6**). Incorporating PEG into the copolymers not only enhanced the colloidal stability of the nanostructures but also increased their biocompatibility. Notably, Zhang and Zhu *et al.*¹⁴⁷ reported water-soluble antibacterial polymer vesicles based on the novel thermo responsive di-block copolymer, which do not contain quaternary ammonium groups (amino groups were protected by Boc and not deprotected) or load any external antibiotics (Fig. 23(b)). Under physiological conditions, these polymer vesicles exhibited superior antibacterial efficacy against both Gram-negative and Gram-positive bacteria. The authors explained that the formation of vesicles increased the local concentration of cationic charges and polymer mass, resulting in stronger interactions between the polymer and the bacterial cell wall/membrane. This led to more effective antibacterial

activity compared to individual, non-self-assembled polymer chains.

Recently, a vesicle capable of real-time visualization of antibacterial processes has been reported in the literature.²⁰⁴ Through modular design, Yang *et al.* integrated multiple functional segments to create vesicles with intrinsic blue fluorescence. The synthesized PCL₂₅-b-PTrp₂-b-P(Lys₁₃-stat-Phe₄) copolymer, featuring biocompatible and biodegradable PCL chains, forms hydrophobic vesicle membranes, while PTrp and P(Lys-stat-Phe) provide intrinsic fluorescence and broad-spectrum antibacterial activity (Fig. 23(c)). Additionally, during the self-assembly process, the aggregation of aromatic amino acids and formation of hydrogen bonds induce electron delocalization, shifting the fluorescence emission from the invisible ultraviolet range of the amino acids to the visible range (with a maximum emission at 436 nm). This transition enables the visualization of the antibacterial process targeting bacteria. Confocal fluorescence imaging of the vesicles with bacteria confirmed the specific adhesion of the vesicles to the bacteria and demonstrated bacterial death resulting from membrane disruption.

In summary, polymer vesicles typically exhibit superior antimicrobial activity compared to their linear counterparts or free antibiotics due to enhanced local charge density or increased delivery efficiency. However, most research on antimicrobial polymer vesicles has focused on those loaded with antibiotics,^{202,205–209} and there is limited work on the design of vesicles incorporating amphiphilic copolymers for antimicrobial purposes. Consequently, this field is still in a phase of rapid growth and discovery, and we believe that many challenges remain to be addressed in the future.



Bacteria strain	Antibacterial agent	Number of passages	MIC (μM)		Fold change in MIC
			Initial	Final	
<i>E. coli</i>	(AB-20) ₁₇	25	12	12	No resistance
	Ciprofloxacin	25	0.188	48.28	256
	(AB-20) ₂₇	30	6.5	6.5	No resistance
	Ciprofloxacin	30	0.09	24.1	256
<i>S. aureus</i>	(AB-20) ₁₇	25	14	14	No resistance
	Ciprofloxacin	25	0.75	48.28	64
	(AB-20) ₂₇	30	6.5	6.5	No resistance
	Ciprofloxacin	30	0.75	24.1	32

The escalating rise in antibiotic resistance among bacteria and other pathogens has emerged as a new threat to global health.²¹⁰ The emergence of this resistance phenomenon is attributed to various factors, including natural evolution under selective pressure, over prescription of antibiotics, and inappropriate use of antibiotics. Due to the development of resistance, there is a growing demand for novel antibiotics. Amphiphilic antimicrobial copolymers exhibit broad-spectrum antibacterial activity and low toxicity, distinct from HDPs, they can physically disrupt the cell membrane of organisms, thereby preventing the development of drug-resistant microorganisms and coupled with inherent advantages in synthetic methods. These attributes have piqued the interest of polymer chemists in exploring the potential of polymeric antibiotics.^{116,211,212}

For example, the amphiphilic antimicrobial copolymers poly[(APMA)-ran-(BMA)], synthesized by A. Tyagi and A. Mishra with approximately 20% hydrophobic benzylmethacrylamide units, were non hemolytic.⁷² Polymers with DP of 17 and 27 exhibited the optimal combination of high antibacterial efficiency and low cytotoxicity. In resistance studies, no increase in MIC values was observed for (AB-20)₁₇ and (AB-20)₂₇ polymers during serial passage over 25 and 30 days, respectively. However, when using ciprofloxacin (a fluoroquinolone antibiotic used as a control), the MIC values for *E. coli* increased by 256-fold, and for *S. aureus*, the MIC values increased by 64-fold (Table 7).

Despite extensive research by both academia and industry, the replacement of antibiotics with amphiphilic antimicrobial copolymers has yet to achieve industrial or clinical application. While these polymers exhibit broad-spectrum antimicrobial activity and a reduced risk of inducing resistance, MICs typically fall in the micromolar range—significantly higher than the nanomolar values of conventional antibiotics.²¹³ This limits their applicability in systemic infection treatment and confines their current use to localized applications such as surface coatings, wound dressings, and medical device modifications.^{17,214} Moreover, balancing hydrophobicity and cationic charge to optimize antimicrobial efficacy without compromising biocompatibility remains a critical challenge. Meanwhile, commercial barriers to the use of amphiphilic antimicrobial copolymers as antibiotic alternatives remain high due to safety concerns, regulatory issues, and challenges in material development.²¹⁵

However, some natural HDPs and their derivatives, such as daptomycin and colistin, have been clinically applied, and several synthetic antimicrobial peptides are currently undergoing clinical trials.²¹⁶ We believe that amphiphilic antimicrobial copolymers will have potential applications as antibiotic alternatives in the future.

4. Conclusion and perspective

In summary, this review systematically summarizes the preparation strategies and wide applications of amphiphilic antibacterial polymers, as well as their advantages in the antibacterial field.

In recent years, scientists have made exciting progress in elucidating the structure–activity relationships of antimicrobial polymers, providing unprecedented tools for the synthesis of amphiphilic antimicrobial copolymers. This has been achieved by chemists through the utilization of actively controlled methods to rigorously control structure, morphology, and topology. For instance, fine-tuning the amphiphilic antimicrobial copolymers balance can enhance selectivity and antimicrobial activity. Currently, the careful design and synthesis of amphiphilic antimicrobial copolymers, aimed at providing a clear understanding of structure/property relationships have become a reality. However, significant work still needs to be undertaken to address major challenges, such as: (1) enhancing antimicrobial activity while mitigating toxicity associated with certain strategies; (2) broadening the range of microbes effectively eliminated without inducing resistance; improving the long-term stability of antimicrobial polymers; (3) developing simple, cost-effective manufacturing solutions; (4) considering the biodegradability of amphiphilic antimicrobial copolymers in specific applications and (5) imbuing amphiphilic antimicrobial copolymers with targeted smart responses.

Moreover, despite the enormous potential applications of amphiphilic antimicrobial copolymers, particularly in biomedical materials such as antibiofilm, wound healing, and tissue engineering, they are still in a rapid growth and discovery phase. Therefore, we believe that there are still many challenges worth attention in the future:

(1) How to design and synthesize amphiphilic antimicrobial copolymers to meet specific application requirements? In certain specific scenarios, antimicrobial agents need to be active against only one or a few microorganisms. This requires a deep understanding of microbial structures and antibacterial mechanisms. Synthesizing such specialized amphiphilic antimicrobial copolymers has consistently been highly challenging.

(2) How to increase the selectivity toward bacteria and mammalian cells? Currently, side effects can be reduced by developing precisely sequence-controlled and highly structured polymers rather than simpler models. Recent studies have found that incorporating neutral hydrophilic groups into traditional binary polymers to create novel ternary antimicrobial polymers can significantly enhance their biocompatibility with host cells. However, our ongoing goal is to further increase

(3) How to achieve antibacterial and anticancer activities simultaneously? In clinical settings, bacterial infections often accompany tumor sites due to the decreased immunity of patients. Therefore, achieving simultaneous antibacterial and anticancer effects in biomedical materials holds special significance. This could bring new insights into the field of cancer treatment.

ABMA	Aminobutyl methacrylate
AEMA	Amino-ethyl methacrylate
AIE	Aggregation-induced emission
AM	4-Acryloylmorpholine
APMA	Aminopropyl methacrylamide
APS	Ammonium persulfate
ATRP	Atom transfer radical polymerization
B	<i>N</i> -Benzylacrylamide
Bm	<i>N</i> -Benzylacrylamide
BMA	Benzylmethacrylamide
CCS	Core-cross-linked star
CPC	Polycarbonate
Cpm	<i>N</i> -Cycloheptylacrylamide
Cxm	<i>N</i> -(Cyclohexylmethyl)acrylamide
DMA	Dopamine methacrylamide
DMC	Methacrylateoethyl trimethyl ammonium chloride
DP	Degree of polymerization
EMA	Ethyl methacrylate
HC ₅₀ and HC ₁₀	The maximum concentration of polymers with less than 50% or 10% hemolytic activity on RBCs
HEA	Hydroxyethyl acrylamide
HEMA	Hydroxyethyl methacrylate
HHMP	2-Hydroxy-4'-(2-hydroxyethoxy)-2-methylpropiophenone
Hm	<i>N</i> -Heptylacrylamide
HPs	Hyperbranched polymers
I	<i>N</i> -Isopentylacrylamide
IC ₅₀	The maximum concentration of polymer less than 50% toxic to test cells
IL	Ionic liquid
Im	<i>N</i> -isopentylacrylamide
G-MPA-N ⁺	Glycidyl methacrylate quaternary ammonium cations
HDPs	Host defense peptides
LPs	Linear polymers
LPS	lipopolysaccharide
L-PTMI	Linear polytrimethyleneimine
MIC	Minimum inhibitory concentration
MPC	Methacryloyloxyethyl phosphorylcholine
MRSA	Methicillin-resistant <i>S. aureus</i>
NCA	<i>N</i> -Carboxyanhydrides
Nm	<i>N</i> -Nonylacrylamide

This is a review article, therefore it does not contain original data.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

The first author of this work was financially supported by the China Scholarship Council (CSC) under grant number 202204910075.

References

- 1 L. J. Rothschild and R. L. Mancinelli, *Nature*, 2001, **409**, 1092–1101.
- 2 G. Antranikian, C. E. Vorgias and C. Bertoldo, *Adv. Biochem. Eng./Biotechnol.*, 2005, **96**, 219–262.
- 3 P. Sarmah, M. M. Dan, D. Adapa and T. Sarangi, *Electron. J. Biol.*, 2018, **14**, 50–58.
- 4 M. Lavanya, *J. Bio-Tribo-Corros.*, 2021, **7**, 125.
- 5 G. McDonnell and A. D. Russell, *Clin. Microbiol. Rev.*, 1999, **12**, 147–179.
- 6 C. Marambio-Jones and E. M. Hoek, *J. Nanopart. Res.*, 2010, **12**, 1531–1551.
- 7 B. K. English and A. H. Gaur, *Hot Topics in Infection and Immunity in Children VI*, 2010, pp. 73–82.
- 8 R. J. Fair and Y. Tor, *Perspect. Med. Chem.*, 2014, **6**, 25–64.
- 9 G. Mancuso, A. Midiri, E. Gerace and C. Biondo, *Pathogens*, 2021, **10**, 1310.
- 10 O. O. Adeniji, M. O. Ojemaye and A. I. Okoh, *ACS Appl. Bio Mater.*, 2022, **5**, 4814–4826.
- 11 R. Rawashdeh and Y. Haik, *Dyn. Biochem. Process Biotechnol. Mol. Biol.*, 2009, **3**, 12–20.
- 12 Y. Yan, Y. Li, Z. Zhang, X. Wang, Y. Niu, S. Zhang, W. Xu and C. Ren, *Colloids Surf., B*, 2021, **202**, 111682.
- 13 J. K. Boparai and P. K. Sharma, *Protein Pept. Lett.*, 2020, **27**, 4–16.
- 14 S. Duan, R. Wu, Y. Xiong, H. Ren, C. Lei, Y. Zhao, X. Zhang and F. Xu, *Prog. Mater. Sci.*, 2022, **125**, 100887.
- 15 K. Huang, C. Yang, S. Huang, C. Chen, Y. Lu and Y. Lin, *Int. J. Mol. Sci.*, 2016, **17**, 1578.
- 16 L. Timofeeva and N. Kleshcheva, *Appl. Microbiol. Biotechnol.*, 2011, **89**, 475–492.
- 17 J. Chen, F. Wang, Q. Liu and J. Du, *Chem. Commun.*, 2014, **50**, 14482–14493.
- 18 A. Jain, L. S. Duvvuri, S. Farah, N. Beyth, A. J. Domb and W. Khan, *Adv. Healthcare Mater.*, 2014, **3**, 1969–1985.
- 19 P. Raffa, D. A. Z. Wever, F. Picchioni and A. A. Broekhuis, *Chem. Rev.*, 2015, **115**, 8504–8563.
- 20 L. C. Klein and A. B. Wojcik, *Encyclopedia of Materials: Science and Technology*, Elsevier, Oxford, 2001, pp. 7577–7584.
- 21 F. Perin, A. Motta and D. Maniglio, *Mater. Sci. Eng., C*, 2021, **123**, 111952.
- 22 A. B. Kutikov and J. Song, *ACS Biomater. Sci. Eng.*, 2015, **1**, 463–480.
- 23 Y. Yang, Z. Cai, Z. Huang, X. Tang and X. Zhang, *Polym. J.*, 2018, **50**, 33–44.
- 24 A. Patrzykat, C. L. Friedrich, L. Zhang, V. Mendoza and R. E. Hancock, *Antimicrob. Agents Chemother.*, 2002, **46**, 605–614.
- 25 A. Krizsan, D. Volke, S. Weinert, N. Sträter, D. Knappe and R. Hoffmann, *Angew. Chem., Int. Ed.*, 2014, **53**, 12236–12239.
- 26 G. Kragol, S. Lovas, G. Varadi, B. A. Condie, R. Hoffmann and L. Otvos, Jr., *Biochemistry*, 2001, **40**, 3016–3026.
- 27 D. G. McCafferty, P. Cudic, M. K. Yu, D. C. Behenna and R. Kruger, *Curr. Opin. Chem. Biol.*, 1999, **3**, 672–680.
- 28 I. Ginsburg, *Med. Hypotheses*, 2004, **62**, 367–374.
- 29 L. Otvos, Jr., H. Flick-Smith, M. Fox, E. Ostorhazi, R. M. Dawson and J. D. Wade, *Protein Pept. Lett.*, 2014, **21**, 374–381.
- 30 Y. Lai and R. L. Gallo, *Trends Immunol.*, 2009, **30**, 131–141.
- 31 D. J. Davidson, A. J. Currie, G. S. Reid, D. M. Bowdish, K. L. MacDonald, R. C. Ma, R. E. Hancock and D. P. Speert, *J. Immunol.*, 2004, **172**, 1146–1156.
- 32 S. V. Guryanova and T. V. Ovchinnikova, *Int. J. Mol. Sci.*, 2022, **23**, 2499.
- 33 H. Li, J. Niu, X. Wang, M. Niu and C. Liao, *Pharmaceutics*, 2023, **15**, 2278.
- 34 L. Brandenburg, J. Merres, L. Albrecht, D. Varoga and T. Pufe, *Polymers*, 2012, **4**, 539–560.
- 35 A. C. Engler, N. Wiradharma, Z. Ong, D. J. Coady, J. L. Hedrick and Y. Yang, *Nano Today*, 2012, **7**, 201–222.
- 36 K. Matsuzaki, *Biochimica et Biophysica Acta*, 2009, **1788**, 1687–1692.
- 37 P. Gilbert and L. E. Moore, *J. Appl. Microbiol.*, 2005, **99**, 703–715.
- 38 M. F. Ilker, K. Nüsslein, G. N. Tew and E. B. Coughlin, *J. Am. Chem. Soc.*, 2004, **126**, 15870–15875.
- 39 H. Mortazavian, L. L. Foster, R. Bhat, S. Patel and K. Kuroda, *Biomacromolecules*, 2018, **19**, 4370–4378.
- 40 G. Moad, E. Rizzardo and S. H. Thang, *Chem. – Asian J.*, 2013, **8**, 1634–1644.
- 41 F. L. Hatton, *Polym. Chem.*, 2020, **11**, 220–229.
- 42 N. P. Truong, G. R. Jones, K. G. Bradford, D. Konkolewicz and A. Anastasaki, *Nat. Rev. Chem.*, 2021, **5**, 859–869.
- 43 N. Corrigan, K. Jung, G. Moad, C. J. Hawker, K. Matyjaszewski and C. Boyer, *Prog. Polym. Sci.*, 2020, **111**, 101311.

- 50 D. S. Uppu, S. Samaddar, C. Ghosh, K. Paramanandham, B. R. Shome and J. Haldar, *Biomaterials*, 2016, **74**, 131–143.
- 51 K. E. Locock, T. D. Michl, J. D. Valentin, K. Vasilev, J. D. Hayball, Y. Qu, A. Traven, H. J. Griesser, L. Meagher and M. Haeussler, *Biomacromolecules*, 2013, **14**, 4021–4031.
- 52 K. Kuroda and W. F. DeGrado, *J. Am. Chem. Soc.*, 2005, **127**, 4128–4129.
- 53 K. Kuroda, G. A. Caputo and W. F. DeGrado, *Chem. – Eur. J.*, 2009, **15**, 1123–1133.
- 54 K. Lienkamp and G. N. Tew, *Chem. – Eur. J.*, 2009, **15**, 11784–11800.
- 55 K. Lienkamp, A. E. Madkour, A. Musante, C. F. Nelson, K. Nusslein and G. N. Tew, *J. Am. Chem. Soc.*, 2008, **130**, 9836–9843.
- 56 G. J. Gabriel, J. A. Maegerlein, C. F. Nelson, J. M. Dabkowski, T. Eren, K. Nüsslein and G. N. Tew, *Chem. – Eur. J.*, 2009, **15**, 433–439.
- 57 B. Dizman, M. O. Elasri and L. J. Mathias, *J. Polym. Sci., Part A: Polym. Chem.*, 2006, **44**, 5965–5973.
- 58 E. R. Kenawy, S. D. Worley and R. Broughton, *Biomacromolecules*, 2007, **8**, 1359–1384.
- 59 T. Tashiro, *Macromol. Mater. Eng.*, 2001, **286**, 63–87.
- 60 J. C. Tiller, C. J. Liao, K. Lewis and A. M. Klibanov, *Proc. Natl. Acad. Sci. U. S. A.*, 2001, **98**, 5981–5985.
- 61 F. Baudrion, A. Perichaud and S. Coen, *J. Appl. Polym. Sci.*, 1998, **70**, 2657–2666.
- 62 E. R. Kenawy, F. I. Abdel-Hay, A. El-Shanshoury and M. H. El-Newehy, *J. Polym. Sci., Part A: Polym. Chem.*, 2002, **40**, 2384–2393.
- 63 M. A. Gelman, B. Weisblum, D. M. Lynn and S. H. Gellman, *Org. Lett.*, 2004, **6**, 557–560.
- 64 N. M. Milovic, J. Wang, K. Lewis and A. M. Klibanov, *Biotechnol. Bioeng.*, 2005, **90**, 715–722.
- 65 R. F. Epand, B. P. Mowery, S. E. Lee, S. S. Stahl, R. I. Lehrer, S. H. Gellman and R. M. Epand, *J. Mol. Biol.*, 2008, **379**, 38–50.
- 66 J. Y. Huang, R. R. Koepsel, H. Murata, W. Wu, S. B. Lee, T. Kowalewski, A. J. Russell and K. Matyjaszewski, *Langmuir*, 2008, **24**, 6785–6795.
- 67 H. Murata, R. R. Koepsel, K. Matyjaszewski and A. J. Russell, *Biomaterials*, 2007, **28**, 4870–4879.
- 68 E. F. Palermo and K. Kuroda, *Biomacromolecules*, 2009, **10**, 1416–1428.
- 69 E. F. Palermo, S. Vemparala and K. Kuroda, *Biomacromolecules*, 2012, **13**, 1632–1641.
- 70 P. Pham, S. Oliver, E. H. Wong and C. Boyer, *Polym. Chem.*, 2021, **12**, 5689–5703.
- 71 X. Yang, K. Hu, G. Hu, D. Shi, Y. Jiang, L. Hui, R. Zhu, Y. Xie and L. Yang, *Biomacromolecules*, 2014, **15**, 3267–3277.
- 72 A. Tyagi and A. Mishra, *ACS Omega*, 2021, **6**, 34724–34735.
- 73 W. Yang, K. G. Neoh, E. T. Kang, S. L. M. Teo and D. Rittschof, *Polym. Chem.*, 2013, **4**, 3105–3115.
- 74 W. Zheng, M. Anzaldúa, A. Arora, Y. Jiang, K. McIntyre, M. Doerfert, T. Winter, A. Mishra, H. Ma and H. Liang, *Biomacromolecules*, 2020, **21**, 2187–2198.
- 75 J. S. Brown, Z. J. Mohamed, C. M. Artim, D. N. Thornlow, J. F. Hassler, V. P. Rigoglioso, S. Daniel and C. A. Alabi, *Commun. Biol.*, 2018, **1**, 220.
- 76 M. Porel, D. N. Thornlow, C. M. Artim and C. A. Alabi, *ACS Chem. Biol.*, 2017, **12**, 715–723.
- 77 P. Pham, S. Oliver and C. Boyer, *Macromol. Chem. Phys.*, 2022, **224**, 2200226.
- 78 A. M. Carmona-Ribeiro and L. D. de Melo Carrasco, *Int. J. Mol. Sci.*, 2013, **14**, 9906–9946.
- 79 Z. Zhang, X. Wang, J. Liu, H. Yang, H. Tang, J. Li, S. Luan, J. Yin, L. Wang and H. Shi, *Angew. Chem., Int. Ed.*, 2024, **63**, e202318011.
- 80 A. Wang, S. Duan, X. Ding, N. Zhao, Y. Hu, X. Ding and F. Xu, *Adv. Funct. Mater.*, 2021, **31**, 2011165.
- 81 H. Takahashi, G. A. Caputo, S. Vemparala and K. Kuroda, *Bioconjugate Chem.*, 2017, **28**, 1340–1350.
- 82 L. Liu, K. Xu, H. Wang, P. K. Tan, W. Fan, S. Venkatraman, L. Li and Y. Yang, *Nat. Nanotechnol.*, 2009, **4**, 457–463.
- 83 A. King, S. Chakrabarty, W. Zhang, X. Zeng, D. E. Ohman, L. F. Wood, S. Abraham, R. Rao and K. J. Wynne, *Biomacromolecules*, 2014, **15**, 456–467.
- 84 I. Mukherjee, A. Ghosh, P. Bhadury and P. De, *ACS Omega*, 2017, **2**, 1633–1644.
- 85 S. Mankoci, R. L. Kaiser, N. Sahai, H. A. Barton and A. Joy, *ACS Biomater. Sci. Eng.*, 2017, **3**, 2588–2597.
- 86 T. Eren, A. Som, J. R. Rennie, C. F. Nelson, Y. Urgina, K. Nüsslein, E. B. Coughlin and G. N. Tew, *Macromol. Chem. Phys.*, 2008, **209**, 516–524.
- 87 Y. Hu, J. Zhao, J. Zhang, Z. Zhu and J. Rao, *ACS Macro Lett.*, 2021, **10**, 990–995.
- 88 J. Oh and A. Khan, *Biomacromolecules*, 2021, **22**, 3534–3542.
- 89 Y. Zhao, X. Wang, Y. Hu, J. Zhao, M. Sun, M. Yang, H. Xuan, X. Wang, J. Zhang, Z. Zhu and J. Rao, *ACS Appl. Polym. Mater.*, 2023, **5**, 4437–4447.
- 90 K. J. Cutrona, B. A. Kaufman, D. M. Figueroa and D. E. Elmore, *FEBS Lett.*, 2015, **589**, 3915–3920.
- 91 F. Nederberg, Y. Zhang, J. P. Tan, K. Xu, H. Wang, C. Yang, S. Gao, X. Guo, K. Fukushima and L. Li, *Nat. Chem.*, 2011, **3**, 409–414.
- 92 L. M. Thoma, B. R. Boles and K. Kuroda, *Biomacromolecules*, 2014, **15**, 2933–2943.
- 93 A. K. Marr, W. J. Gooderham and R. E. Hancock, *Curr. Opin. Pharmacol.*, 2006, **6**, 468–472.
- 94 G. Maisetta, M. Di Luca, S. Esin, W. Florio, F. L. Brancatisano, D. Bottai, M. Campa and G. Batoni, *Peptides*, 2008, **29**, 1–6.
- 95 S. Hong, H. Takahashi, E. T. Nadres, H. Mortazavian, G. A. Caputo, J. G. Younger and K. Kuroda, *PLoS One*, 2017, **12**, e0169262.
- 96 R. Bhat, L. L. Foster, G. Rani, S. Vemparala and K. Kuroda, *RSC Adv.*, 2021, **11**, 22044–22056.
- 97 S. Buffet-Bataillon, P. Tattevin, M. Bonnaure-Mallet and A. Jolivet-Gougeon, *Int. J. Antimicrob. Agents*, 2012, **39**, 381–389.
- 98 S. Liu, R. J. Ono, H. Wu, J. Y. Teo, Z. C. Liang, K. Xu, M. Zhang, G. Zhong, J. P. Tan, M. Ng, C. Yang, J. Chan,

Z. Ji, C. Bao, K. Rao, S. Kumar, S. Gao, A. Lee, M. Fevre, H. Dong, J. Y. Ying, L. Li, W. Fan, J. L. Hedrick and Y. Y. Yang, *Biomaterials*, 2017, **127**, 36–48.

99 W. Chin, G. Zhong, Q. Pu, C. Yang, W. Lou, P. F. De Sessions, B. Periaswamy, A. Lee, Z. Liang, X. Ding, S. Gao, C. W. Chu, S. Bianco, C. Bao, Y. Tong, W. Fan, M. Wu, J. Hedrick and Y. Yang, *Nat. Commun.*, 2018, **9**, 917.

100 J. Leong, C. Yang, J. Tan, B. Tan, S. Hor, J. L. Hedrick and Y. Yang, *Biomater. Sci.*, 2020, **8**, 6920–6929.

101 J. S. Brown, Z. J. Mohamed, C. M. Artim, D. N. Thornlow, J. F. Hassler, V. P. Rigoglioso, S. Daniel and C. A. Alabi, *Commun. Biol.*, 2018, **1**, 220.

102 V. Sambhy, B. R. Peterson and A. Sen, *Angew. Chem., Int. Ed.*, 2008, **47**, 1250–1254.

103 M. N. Islam, B. Aksu, M. Güncü, M. Gallei, M. Tulu and T. Eren, *J. Polym. Sci.*, 2020, **58**, 872–884.

104 R. Cuervo-Rodríguez, A. Muñoz-Bonilla, F. López-Fabal and M. Fernández-García, *Polymers*, 2020, **12**, 972.

105 S. N. Hancock, N. Yuntawattana, E. Diep, A. Maity, A. Tran, J. D. Schiffman and Q. Michaudel, *Proc. Natl. Acad. Sci. U. S. A.*, 2023, **120**, e2311396120.

106 A. Vishwakarma, F. Dang, A. Ferrell, H. A. Barton and A. Joy, *J. Am. Chem. Soc.*, 2021, **143**, 9440–9449.

107 Z. Zheng, Q. Xu, J. Guo, J. Qin, H. Mao, B. Wang and F. Yan, *ACS Appl. Mater. Interfaces*, 2016, **8**, 12684–12692.

108 Z. Shi, X. Zhang, Z. Yu, F. Yang, H. Liu, R. Xue, S. Luan and H. Tang, *Biomacromolecules*, 2021, **22**, 2373–2381.

109 A. Kanazawa, T. Ikeda and T. Endo, *J. Polym. Sci., Part A: Polym. Chem.*, 1993, **31**, 335–343.

110 B. Hisey, P. J. Ragona and E. R. Gillies, *Biomacromolecules*, 2017, **18**, 914–923.

111 H. Kудay, N. C. Süer, A. Bayır, B. Aksu, A. Hatipoğlu, M. M. Güncü, İ. Acaroğlu Degitz, M. Gallei and T. Eren, *ACS Appl. Polym. Mater.*, 2021, **3**, 6524–6538.

112 A. Kanazawa, T. Ikeda and T. Endo, *J. Polym. Sci., Part A: Polym. Chem.*, 1993, **31**, 2873–2876.

113 P. T. Phuong, S. Oliver, J. He, E. H. H. Wong, R. T. Mathers and C. Boyer, *Biomacromolecules*, 2020, **21**, 5241–5255.

114 E. F. Palermo and K. Kuroda, *Appl. Microbiol. Biotechnol.*, 2010, **87**, 1605–1615.

115 C. Ergene, K. Yasuhara and E. F. Palermo, *Polym. Chem.*, 2018, **9**, 2407–2427.

116 H. Takahashi, G. A. Caputo and K. Kuroda, *Biomater. Sci.*, 2021, **9**, 2758–2767.

117 E. F. Palermo, I. Sovadinova and K. Kuroda, *Biomacromolecules*, 2009, **10**, 3098–3107.

118 I. Sovadinova, E. F. Palermo, M. Urban, P. Mpiğa, G. A. Caputo and K. Kuroda, *Polymers*, 2011, **3**, 1512–1532.

119 M. Ward, M. Sanchez, M. O. Elasri and A. B. Lowe, *J. Appl. Polym. Sci.*, 2006, **101**, 1036–1041.

120 J. Tan, Y. Zhao, J. L. Hedrick and Y. Yang, *Adv. Healthcare Mater.*, 2022, **11**, 2100482.

121 W. Chin, G. Zhong, Q. Pu, C. Yang, W. Lou, P. F. De Sessions, B. Periaswamy, A. Lee, Z. C. Liang, X. Ding, S. Gao, C. W. Chu, S. Bianco, C. Bao, Y. W. Tong, W. Fan, M. Wu, J. L. Hedrick and Y. Y. Yang, *Nat. Commun.*, 2018, **9**, 917.

122 C. Peng, A. Vishwakarma, S. Mankoci, H. A. Barton and A. Joy, *Biomacromolecules*, 2019, **20**, 1675–1682.

123 S. Mankoci, J. Ewing, P. Dalai, N. Sahai, H. A. Barton and A. Joy, *Biomacromolecules*, 2019, **20**, 4096–4106.

124 M. A. Rahman, M. Bam, E. Luat, M. S. Jui, M. S. Ganewatta, T. Shokfai, M. Nagarkatti, A. W. Decho and C. Tang, *Nat. Commun.*, 2018, **9**, 5231.

125 B. C. Allison, B. M. Applegate and J. P. Youngblood, *Biomacromolecules*, 2007, **8**, 2995–2999.

126 R. Namivandi-Zangeneh, R. J. Kwan, T.-K. Nguyen, J. Yeow, F. L. Byrne, S. H. Oehlers, E. H. Wong and C. Boyer, *Polym. Chem.*, 2018, **9**, 1735–1744.

127 C. Ergene and E. F. Palermo, *J. Mater. Chem. B*, 2018, **6**, 7217–7229.

128 S. Schaefer, D. Melodia, C. Pracey, N. Corrigan, M. D. Lenardon and C. Boyer, *Biomacromolecules*, 2024, **25**, 871–889.

129 P. Pham, S. Oliver, D. T. Nguyen and C. Boyer, *Macromol. Rapid Commun.*, 2022, **43**, 2200377.

130 E. T. Nadres, H. Takahashi and K. Kuroda, *J. Polym. Sci., Part A: Polym. Chem.*, 2016, **55**, 304–312.

131 T. D. Michl, K. E. S. Locock, N. E. Stevens, J. D. Hayball, K. Vasilev, A. Postma, Y. Qu, A. Traven, M. Haeussler, L. Meagher and H. J. Griesser, *Polym. Chem.*, 2014, **5**, 5813–5822.

132 W. Chin, C. Yang, V. W. L. Ng, Y. Huang, J. Cheng, Y. W. Tong, D. J. Coady, W. Fan, J. L. Hedrick and Y. Y. Yang, *Macromolecules*, 2013, **46**, 8797–8807.

133 K. Lienkamp, K. N. Kumar, A. Som, K. Nüsslein and G. N. Tew, *Chem. – Eur. J.*, 2009, **15**, 11710–11714.

134 J. Pachla, R. J. Kopiasz, G. Marek, W. Tomaszewski, A. Głogowska, K. Drężek, S. Kowalczyk, R. Podgórski, B. Butruk-Raszeja, T. Ciach, J. Mierzejewska, A. Plichta, E. Augustynowicz-Kopeć and D. Jańczewski, *Biomacromolecules*, 2023, **24**, 2237–2249.

135 Y. Qiao, C. Yang, D. J. Coady, Z. Ong, J. L. Hedrick and Y. Yang, *Biomaterials*, 2012, **33**, 1146–1153.

136 S. Oliver, L. Zhao, A. J. Gormley, R. Chapman and C. Boyer, *Macromolecules*, 2019, **52**, 3–23.

137 D. A. Tomalia, *Prog. Polym. Sci.*, 2005, **30**, 294–324.

138 C. Boyer, V. Bulmus, T. P. Davis, V. Ladmirał, J. Liu and S. Perrier, *Chem. Rev.*, 2009, **109**, 5402–5436.

139 B. D. Fairbanks, P. A. Gunatillake and L. Meagher, *Adv. Drug Delivery Rev.*, 2015, **91**, 141–152.

140 H. Takahashi, E. T. Nadres and K. Kuroda, *Biomacromolecules*, 2017, **18**, 257–265.

141 S. Zhao, W. Huang, C. Wang, Y. Wang, Y. Zhang, Z. Ye, J. Zhang, L. Deng and A. Dong, *Biomacromolecules*, 2020, **21**, 5269–5281.

142 Y. Wang, J. Xu, Y. Zhang, H. Yan and K. Liu, *Macromol. Biosci.*, 2011, **11**, 1499–1504.

143 A. Kuroki, P. Sangwan, Y. Qu, R. Peltier, C. Sanchez-Cano, J. Moat, C. G. Dowson, E. G. Williams, K. E. Locock and M. Hartlieb, *ACS Appl. Mater. Interfaces*, 2017, **9**, 40117–40126.

144 P. R. Judzewitsch, T.-K. Nguyen, S. Shanmugam, E. H. H. Wong and C. Boyer, *Angew. Chem., Int. Ed.*, 2018, **57**, 4559–4564.

- 145 H. Takahashi, I. Sovadinova, K. Yasuhara, S. Vemparala, G. A. Caputo and K. Kuroda, *Wiley Interdiscip. Rev.: Nanomed. Nanobiotechnol.*, 2023, **15**, e1866.
- 146 K. Zhang, Y. Du, Z. Si, Y. Liu, M. E. Turvey, C. Raju, D. Keogh, L. Ruan, S. L. Jothy and S. Reghu, *Nat. Commun.*, 2019, **10**, 4792.
- 147 C. Zhang, Y. Zhu, C. Zhou, W. Yuan and J. Du, *Polym. Chem.*, 2013, **4**, 255–259.
- 148 W. Yuan, J. Wei, H. Lu, L. Fan and J. Du, *Chem. Commun.*, 2012, **48**, 6857–6859.
- 149 C. Chang, C. Chang, Y. Yang, H. Chen, S. Yang, W. Yao and C. Chao, *Polymers*, 2022, **14**, 250.
- 150 D. J. Siegwart, J. K. Oh and K. Matyjaszewski, *Prog. Polym. Sci.*, 2012, **37**, 18–37.
- 151 Y. Oda, S. Kanaoka, T. Sato, S. Aoshima and K. J. B. Kuroda, *Biomacromolecules*, 2011, **12**, 3581–3591.
- 152 Y. Oda, K. Yasuhara, S. Kanaoka, T. Sato, S. Aoshima and K. Kuroda, *Polymers*, 2018, **10**, 93.
- 153 M. Casanova, H. Olleik, S. Hdouech, C. Roblin, J.-F. Cavalier, V. Point, K. Jeannot, B. Caron, J. Perrier, S. Charriau, M. Lafond, Y. Guillaneuf, S. Canaan, C. Lefay and M. Maresca, *Antibiotics*, 2023, **12**, 120.
- 154 Y. Tominaga, M. Mizuse, A. Hashidzume, Y. Morishima and T. Sato, *J. Phys. Chem. B*, 2010, **114**, 11403–11408.
- 155 C. Chen and T. Weil, *Nanoscale Horiz.*, 2022, **7**, 1121–1135.
- 156 J. Koehbach and D. J. Craik, *Trends Pharmacol. Sci.*, 2019, **40**, 517–528.
- 157 J. Xu, L. Pu, J. Ma, S. K. Kumar and H. Duan, *Polym. Chem.*, 2020, **11**, 6632–6639.
- 158 M. Aquib, W. Yang, L. Yu, V. K. Kannaujiya, Y. Zhang, P. Li, A. Whittaker, C. Fu and C. Boyer, *Chem. Sci.*, 2024, **15**, 19057–19069.
- 159 R. Liénard, J. De Winter and O. Coulembier, *J. Polym. Sci.*, 2020, **58**, 1481–1502.
- 160 F. M. Haque and S. M. Grayson, *Nat. Chem.*, 2020, **12**, 433–444.
- 161 M. R. E. Santos, P. V. Mendonça, M. C. Almeida, R. Branco, A. C. Serra, P. V. Morais and J. F. J. Coelho, *Biomacromolecules*, 2019, **20**, 1146–1156.
- 162 M. Aquib, S. Schaefer, H. Gmedhin, N. Corrigan, V. A. Bobrin and C. Boyer, *Eur. Polym. J.*, 2024, **205**, 112698.
- 163 S. Laroque, K. E. S. Locock and S. Perrier, *Biomacromolecules*, 2024, **26**, 190–200.
- 164 K. A. Gibney, I. Sovadinova, A. I. Lopez, M. Urban, Z. Ridgway, G. A. Caputo and K. Kuroda, *Macromol. Biosci.*, 2012, **12**, 1279–1289.
- 165 H. Liu, X. Zhang, Z. Zhao, F. Yang, R. Xue, L. Yin, Z. Song, J. Cheng, S. Luan and H. Tang, *Biomater. Sci.*, 2021, **9**, 2721–2731.
- 166 N. Hadjesfandiari, K. Yu, Y. Mei and J. N. Kizhakkedathu, *J. Mater. Chem. B*, 2014, **2**, 4968–4978.
- 167 T. Chen, H. Yang, X. Wu, D. Yu, A. Ma, X. He, K. Sun and J. Wang, *Langmuir*, 2019, **35**, 3031–3037.
- 168 H. Luo, X. Q. Yin, P. F. Tan, Z. P. Gu, Z. M. Liu and L. Tan, *J. Mater. Chem. B*, 2021, **9**, 2802–2815.
- 169 M. R. E. Santos, A. C. Fonseca, P. V. Mendonça, R. Branco, A. C. Serra, P. V. Morais and J. F. J. Coelho, *Materials*, 2016, **9**, 599.
- 170 M. M. Konai, B. Bhattacharjee, S. Ghosh and J. Haldar, *Biomacromolecules*, 2018, **19**, 1888–1917.
- 171 M. Cloutier, D. Mantovani and F. Rosei, *Trends Biotechnol.*, 2015, **33**, 637–652.
- 172 S. Arora, V. Yadav, P. Kumar, R. Gupta and D. Kumar, *Int. J. Pharm. Sci. Rev. Res.*, 2013, **23**, 279–290.
- 173 J. Hoque, S. Ghosh, K. Paramanandham and J. Haldar, *ACS Appl. Mater. Interfaces*, 2019, **11**, 39150–39162.
- 174 Z. Ma, Y. Liu, K. Feng, J. Wei, J. Liu, Y. Wu, X. Pei, B. Yu, M. Cai and F. Zhou, *ACS Appl. Mater. Interfaces*, 2022, **14**, 18901–18909.
- 175 S. Bai, X. Li, Y. Zhao, L. Ren and X. Yuan, *ACS Appl. Mater. Interfaces*, 2020, **12**, 12305–12316.
- 176 C. Chen, Z. Li, X. Li, C. Kuang, X. Liu, Z. Song, H. Liu and Y. Shan, *Composites, Part B*, 2022, **232**, 109623.
- 177 W. E. Hennink and C. F. van Nostrum, *Adv. Drug Delivery Rev.*, 2012, **64**, 223–236.
- 178 H. Chang, C. Li, R. Huang, R. Su, W. Qi and Z. He, *J. Mater. Chem. B*, 2019, **7**, 2899–2910.
- 179 H. Qu, Q. Yao, T. Chen, H. Wu, Y. Liu, C. Wang and A. Dong, *Adv. Colloid Interface Sci.*, 2024, **325**, 103099.
- 180 C. M. González-Henríquez, M. A. Sarabia-Vallejos and J. Rodríguez-Hernández, *Materials*, 2017, **10**, 232.
- 181 M. Malmsten, *Soft Matter*, 2011, **7**, 8725–8736.
- 182 X. Qin, K. Labuda, J. Chen, V. Hruschka, A. Khadem, R. Liska, H. Redl and P. Slezak, *Adv. Funct. Mater.*, 2015, **25**, 6606–6617.
- 183 X. Zhao, H. Wu, B. Guo, R. Dong, Y. Qiu and P. X. Ma, *Biomaterials*, 2017, **122**, 34–47.
- 184 Y. Li, K. Fukushima, D. J. Coady, A. C. Engler, S. Liu, Y. Huang, J. S. Cho, Y. Guo, L. S. Miller and J. P. Tan, *Angew. Chem., Int. Ed.*, 2013, **52**, 674–678.
- 185 N. Zhao and W. Yuan, *Composites, Part B*, 2022, **230**, 109525.
- 186 Y. Hong, Y. Xi, J. Zhang, D. Wang, H. Zhang, N. Yan, S. He and J. Du, *J. Mater. Chem. B*, 2018, **6**, 6311–6321.
- 187 Y. Wang, J. Zhao, Z. Dong, C. Wang, H. Meng, Y. Li, H. Jin and C. Wang, *Mater. Today Chem.*, 2021, **21**, 100537.
- 188 S. Brahmachari, S. Debnath, S. Dutta and P. K. Das, *Beilstein J. Org. Chem.*, 2010, **6**, 859–868.
- 189 S. Li, S. Dong, W. Xu, S. Tu, L. Yan, C. Zhao, J. Ding and X. Chen, *Adv. Sci.*, 2018, **5**, 1700527.
- 190 S. Vidovic, J. Stojkovska, M. Stevanovic, B. Balanc, M. Vukasinovic-Sekulic, A. Marinkovic and B. Obradovic, *R. Soc. Open Sci.*, 2022, **9**, 211517.
- 191 K. Neibert, V. Gopishetty, A. Grigoryev, I. Tokarev, N. Al-Hajaj, J. Vorstenbosch, A. Philip, S. Minko and D. Maysinger, *Adv. Healthcare Mater.*, 2012, **1**, 621–630.
- 192 C. T. Gustafson, F. Boakye-Agyeman, C. L. Brinkman, J. M. Reid, R. Patel, Z. Bajzer, M. Dadsetan and M. J. Yaszemski, *PLoS One*, 2016, **11**, e0146401.
- 193 S. C. Bondy, *Curr. Opin. Toxicol.*, 2021, **26**, 8–13.
- 194 K. Gwon, I. Han, S. Lee, Y. Kim and D. N. Lee, *ACS Appl. Mater. Interfaces*, 2020, **12**, 20234–20242.
- 195 D. Han, Y. Li, X. Liu, B. Li, Y. Han, Y. Zheng, K. W. K. Yeung, C. Li, Z. Cui, Y. Liang, Z. Li, S. Zhu, X. Wang and S. Wu, *Chem. Eng. J.*, 2020, **396**, 125194.

- 196 B. Zeng, Z. Cai, J. Lalevée, Q. Yang, H. Lai, P. Xiao, J. Liu and F. Xing, *Toxicol. In Vitro*, 2021, **72**, 105103.
- 197 D. E. Discher and A. Eisenberg, *Science*, 2002, **297**, 967–973.
- 198 M. A. Rahman, M. S. Jui, M. Bam, Y. Cha, E. Luat, A. Alabresm, M. Nagarkatti, A. W. Decho and C. Tang, *ACS Appl. Mater. Interfaces*, 2020, **12**, 21221–21230.
- 199 Z. Shi, K. G. Neoh, E. T. Kang and W. Wang, *Biomaterials*, 2006, **27**, 2440–2449.
- 200 H. Zhu, Q. Geng, W. Chen, Y. Zhu, J. Chen and J. Du, *J. Mater. Chem. B*, 2013, **1**, 5496–5504.
- 201 K. Lienkamp, K. N. Kumar, A. Som, K. Nüsslein and G. N. Tew, *Chemistry*, 2009, **15**, 11710–11714.
- 202 L. D. Blackman, Z. Y. Oo, Y. Qu, P. A. Gunatillake, P. Cass and K. E. S. Locock, *ACS Appl. Mater. Interfaces*, 2020, **12**, 11353–11362.
- 203 N. J. Bassous and T. J. Webster, *Small*, 2019, **15**, 1804247.
- 204 Y. Yang, L. Chen, M. Sun, C. Wang, Z. Fan and J. Du, *Chin. J. Polym. Sci.*, 2021, **39**, 1412–1420.
- 205 Y. Zhu, B. Yang, S. Chen and J. Du, *Prog. Polym. Sci.*, 2017, **64**, 1–22.
- 206 H. Sun, Y. Wang and J. Song, *Polymers*, 2021, **13**, 2903.
- 207 Y. Chen, Y. Huang and Q. Jin, *Macromol. Chem. Phys.*, 2022, **223**, 2100440.
- 208 T. Wang, Y. Li, E. J. Cornel, C. Li and J. Du, *ACS Nano*, 2021, **15**, 9027–9038.
- 209 Y. Li, G. Liu, X. Wang, J. Hu and S. Liu, *Angew. Chem., Int. Ed.*, 2016, **55**, 1760–1764.
- 210 C. J. Murray, K. S. Ikuta, F. Sharara, L. Swetschinski, G. R. Aguilar, A. Gray, C. Han, C. Bisignano, P. Rao and E. Wool, *Lancet*, 2022, **399**, 629–655.
- 211 S. Alfei and A. M. Schito, *Polymers*, 2020, **12**, 1195.
- 212 H. Etayash and R. E. W. Hancock, *Pharmaceutics*, 2021, **13**, 1820.
- 213 N. F. Kamaruzzaman, L. P. Tan, R. H. Hamdan, S. S. Choong, W. K. Wong, A. J. Gibson, A. Chivu and M. d F. Pina, *Int. J. Mol. Sci.*, 2019, **20**, 2747.
- 214 K. Malhotra and Y. Singh, *Racing for the Surface: Pathogenesis of Implant Infection and Advanced Antimicrobial Strategies*, 2020, pp. 543–581.
- 215 R. Namivandi-Zangeneh, E. H. H. Wong and C. Boyer, *ACS Infect. Dis.*, 2021, **7**, 215–253.
- 216 C. H. Chen and T. K. Lu, *Antibiotics*, 2020, **9**, 24.

