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Conversion of oligo(ethyleneglycol)s into non-toxic highly selective biocompatible poly(ethyleneglycol)s: synthesis, antimicrobial and antibiofilm activity†

Sulbha Kumari,^{‡a} Arpita Halder,^{‡b} Aayush Anand,^a Oindrilla Mukherjee^{*b} and Subrata Chattopadhyay ^{*a}

Designing non-toxic, non-hemolytic, selective antimicrobials remains an important and challenging research problem. Herein, we report an affordable synthetic route to prepare a series of ten multifunctional polyethylene glycols (PEGs) via a cascade reaction approach involving aza-Michael polyaddition followed by post-polymerization modifications using triazolinedione-based click reactions. All polymers are characterized by NMR, IR, SEC, DSC and TG analyses. Antimicrobial and hemolytic studies reveal that structure plays a pivotal role in tuning the antimicrobial efficacy and selectivity (HC/MIC) of the functional PEGs. The selectivity (HC/MIC) reported for the best prototype (InPEG₇₀₀-C₁₂-TAD) is 129, 33 and 39 against *P. aeruginosa*, *E. coli* and *S. aureus*, respectively. Additionally, all the polymers are non-cytotoxic, as revealed by the MTT assay, and exhibit excellent antibiofilm activity.

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1. Introduction

Infectious diseases are increasingly becoming a global problem in today's society, making the suitable design of antibiotics and antimicrobials a very important research topic.^{1–3} Since the first antibiotic 'penicillin' was reported by Alexander Fleming,⁴ several other antibiotics (low molecular weight and polymeric) have been reported.⁵ However, most conventional antibiotics are becoming increasingly ineffective because of antibiotic resistance, which results in multi-drug-resistant microbes. On the other hand, host defence peptides or antimicrobial peptides, which are an invariable part of many living organisms, defend host organisms against microbial infections effectively. Most of these antimicrobial peptides are composed of polymer chains of amino acids that include several cationic amino and hydrophobic functionalities. Due to their cationic nature, these peptides kill negatively charged bacteria via contact killing-membrane disruption mechanisms, which are explained via different models in the literature

including the barrel-stave model, carpet-like model, toroidal-pore model and disordered toroidal-pore model.^{6–10} Due to the contact-active membrane disruption mechanism, such antimicrobial peptides are less able to develop antimicrobial resistance. However, it is extremely challenging to develop such antimicrobial peptides to combat infections as part of our day-to-day needs due to their lower synthetic upscalability, higher cost and the requirement to further improve their selectivity, indicated by the balance between antimicrobial and hemolytic activity.^{11–13} To address these issues, synthesis of contact-active antimicrobial polymers (or polymers showing intrinsic antimicrobial activity), which mimic the structural features of antimicrobial peptides, such as polymeric structure and the tuneable presence of cationic and hydrophobic groups within the backbone, has emerged as a very important research topic.^{14–16} In the literature, various antimicrobial polymers with diverse functional backbones have been reported, including polyacrylates,^{17–19} polyacrylamide,^{20,21} polystyrene,²² polyethyleneamine,²³ polyvinylamine,^{24,25} chitosan,^{26–29} polytetrahydrofurans,³⁰ polyoxazolines,^{31,32} polysulphoniums,^{33,34} etc. Although antimicrobial properties can be tuned via adjusting the polymer's structure (hydrophobic/cationic balance), many of them suffer from higher hemolytic activity, resulting in lower selectivity (MIC/HC₅₀), low biocompatibility and poor water solubility.

Herein, we aim to synthesize poly(ethyleneglycol) (PEG)-based contact-active antimicrobial polymers. PEGs are well known for their biocompatibility, non-immunogenicity and

^aDepartment of Chemistry, Indian Institute of Technology Patna, Bihta, Patna 801106, Bihar, India. E-mail: sch@iitp.ac.in

^bDepartment of Biotechnology, National Institute of Technology, Durgapur, Mahatma Gandhi Rd, A-Zone, Durgapur, West Bengal 713209, India. E-mail: omukherjee.bt@nitdgp.ac.in

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‡These authors contributed equally to the current work.



reactions, enabling the further introduction of various alkyl chains to fine-tune both antimicrobial activity and selectivity (HC_{50}/MIC).

2. Results and discussion

2.1 Synthesis of an indole-functionalized poly(ethylene glycol) polymer (InPEG_x) and its post-polymerization modifications (InPEG_x-C_n-TAD)

A versatile indole-functionalized polyethylene glycol (PEG) periodic copolymer, InPEG_x, was designed and synthesized such

that the biocompatible PEG backbone enhances its aqueous solubility and biocompatibility, while the indole and amine moieties confer antimicrobial activity. As illustrated in Scheme 1 (step 1), polymerization proceeds *via* an aza-Michael polyaddition reaction between commercially available PEG diacrylates (two analogues with $M_n = 575$ and 700 used as macromonomers) and tryptamine in a 1 : 1 molar ratio using a 2 : 1 (v/v) mixture of water and DMF. The two PEG copolymers have different periodic nature, which was modulated by changing the PEG chain length (InPEG_x; $x = 575$ and 700). The synthesized polymers were characterized using ¹H- and ¹³C-NMR, HSQC and SEC analyses.



Fig. 2 (A) ¹H-NMR spectra and (B) ¹³C-NMR spectra of InPEG₅₇₅ and PEG diacrylate (macromonomer).



The ^1H -NMR spectrum of InPEG₅₇₅ (Fig. 2A) clearly revealed the successful insertion of the indole functionality into the PEG copolymer, as evidenced by the appearance of characteristic aromatic and NH peaks of indole functionality between 6.9 and 7.6 ppm and at 10.6 ppm, respectively, along with the characteristic peaks for PEG. The ^{13}C -NMR spectra (Fig. 2B) also provide strong evidence for the insertion of indole *via* an aza-Michael reaction. The characteristic methylene peaks at 23 ppm ($-\text{NH}-\text{CH}_2-\text{CH}_2-\text{Ar}$) and 49 ppm ($-\text{NH}-\text{CH}_2-\text{CH}_2-\text{Ar}$) along with aromatic peaks at 111.8–127.7 ppm and 136.7 ppm are consistent with the indole moiety. Furthermore, the appearance of new methylene peaks at 32.5 ppm ($\text{OCO}-\text{CH}_2-\text{CH}_2-\text{N}$) and 54.3 ppm ($\text{OCO}-\text{CH}_2-\text{CH}_2-\text{N}$) confirms the successful aza-Michael reaction between the acrylate (end) functionality and the amine group. HSQC spectra of InPEG₅₇₅ (Fig. S1†) supported the same molecular details. Similar NMR evidence was observed for InPEG₇₀₀ (Fig. S2†). The SEC analysis revealed the M_n of 5300 and 6100 g mol^{-1} for InPEG₅₇₅ and InPEG₇₀₀ copolymers, respectively.

In the next stage, the indole group serves as an effective clickable site for triazolinedione (TAD) molecules *via* an ultra-fast TAD–indole reaction under ambient reaction conditions.^{39–42} This facile approach offers a highly efficient method for introducing hydrophobic groups as pendant functionalities within polymer chains. For the purpose of post-polymerization modification, four different triazolinedione derivatives, having different alkyl chain lengths (butyl-, hexyl-, octyl- and dodecyl-TAD), were synthesized using a previously reported procedure and characterized using NMR, FTIR and mass spectroscopy.^{43,44}

The four synthesized TAD derivatives with varying alkyl chain lengths (C₄-TAD, C₆-TAD, C₈-TAD and C₁₂-TAD) were subsequently employed for post-modification of InPEG_x *via* a TAD–indole click reaction. The TAD–indole click reaction was initiated by combining solutions of TAD and InPEG_x in DMF at ambient temperature. After a 20 minute reaction period, the

solvent was removed *via* lyophilization. Using this generalized approach, eight sets of hydrophobic post-modified PEG copolymers, with varying alkyl chains and PEG chain lengths, were prepared (InPEG_x-C_n-TAD; $x = 575$ and 700 , $n = 4, 6, 8, 12$), and all the modified copolymers were characterized using NMR, FTIR and SEC analyses. From the ^1H -NMR spectra of the post-modified copolymer InPEG₅₇₅-C₄-TAD (Fig. 3A), the appearance of new characteristic peaks for the butyl group of the TAD moiety at 0.8, 1.0–1.55 and 3.4 ppm provided evidence of alkyl functionalization within the polymer backbone. Furthermore, the TAD–indole adduct was confirmed using ^{13}C -NMR spectra (Fig. 3A). The presence of characteristic butyl peaks at 13.9, 19.7, 30.0 and 38.0 ppm, along with the appearance of a new peak at ~ 77.9 ppm corresponding to the quaternary carbon after the TAD–indole linkage, a peak at ~ 155.6 ppm associated with the carbonyl carbon of the TAD unit, and a peak at 163.8 ppm that corresponds to $\text{N}=\text{CH}$ of the indole unit supports the successful conjugation of the TAD moiety to the indole group of the copolymer. A similar observation was made in the NMR spectra of other hydrophobic TAD-modified InPEG₅₇₅ and InPEG₇₀₀ polymers (Fig. S3–S6†). Subsequently, SEC measurements were performed, and the resulting molecular weights, summarized in Table 1, support the successful post-polymerization modifications.

Table 1 Details of the molecular weights of the unmodified and modified InPEG_x polymers obtained from SEC measurements

Sample	M_n (g mol^{-1})	D	Sample	M_n (g mol^{-1})	D
InPEG ₅₇₅	5300	1.4	InPEG ₇₀₀	6100	1.3
InPEG ₅₇₅ -C ₄ -TAD	5800	1.4	InPEG ₇₀₀ -C ₄ -TAD	6200	1.4
InPEG ₅₇₅ -C ₆ -TAD	5830	1.4	InPEG ₇₀₀ -C ₆ -TAD	6350	1.4
InPEG ₅₇₅ -C ₈ -TAD	6200	1.3	InPEG ₇₀₀ -C ₈ -TAD	6630	1.4
InPEG ₅₇₅ -C ₁₂ -TAD	6280	1.3	InPEG ₇₀₀ -C ₁₂ -TAD	6360	1.3



Fig. 3 (A) ^1H - and ^{13}C -NMR spectra of InPEG₅₇₅-C₄-TAD and (B) FTIR spectra of InPEG₅₇₅ and InPEG₅₇₅-C_n-TAD ($n = 4, 6, 8, 12$).



Furthermore, the synthesis of indolized PEGs and their post-polymer modifications were also evidenced by FTIR spectral analysis. Fig. 3B depicts the FTIR spectra of un-

modified and modified InPEG₅₇₅ with various mono-TADs. In the case of unmodified InPEG₅₇₅, the appearance of bands at 3055 and 3364 cm⁻¹, corresponding to the aro-



Fig. 4 TGA thermograms of (A) InPEG₅₇₅ and its four post-functional derivatives and (B) InPEG₇₀₀ and its post-functional derivatives.

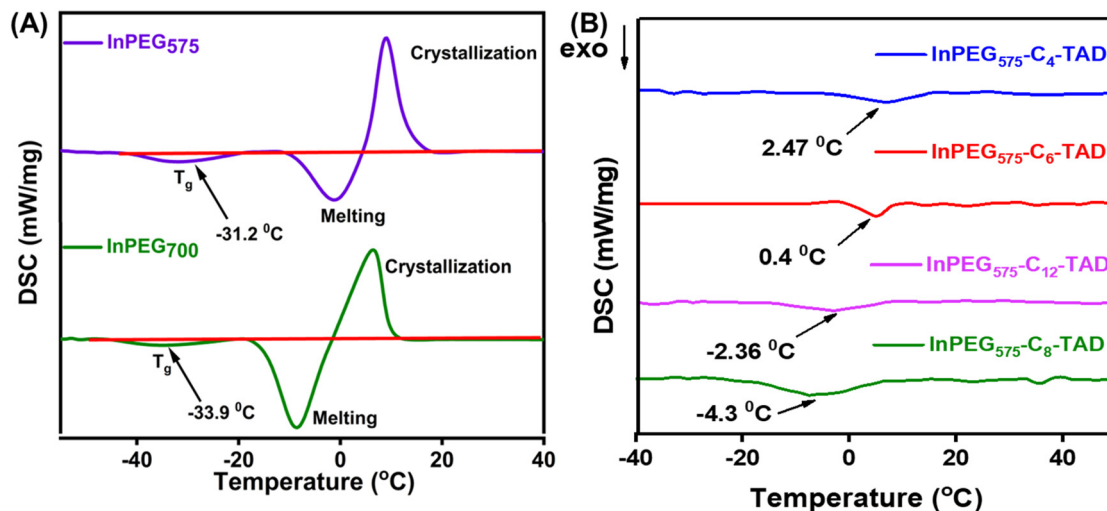


Fig. 5 DSC thermograms of (A) InPEG₅₇₅ and InPEG₇₀₀ and (B) post-polymerization derivatives of InPEG₅₇₅-C_n-TAD.

Table 2 Antimicrobial activity and hemolytic assay results of the polymers

Polymer	MIC ($\mu\text{g mL}^{-1}$)			HC ₅₀ ($\mu\text{g mL}^{-1}$)	Selectivity (HC/MIC)		
	<i>P. aeruginosa</i>	<i>E. coli</i>	<i>S. aureus</i>		<i>P. aeruginosa</i>	<i>E. coli</i>	<i>S. aureus</i>
InPEG ₅₇₅	162	587	385	7130	44.13	12.14	18.50
InPEG ₅₇₅ -C ₄ -TAD	211	579	401	6582	31.19	11.36	16.43
InPEG ₅₇₅ -C ₆ -TAD	113	526	269	6760	59.93	12.84	25.15
InPEG ₅₇₅ -C ₈ -TAD	44	160	156	2022	46.42	12.66	13.00
InPEG ₅₇₅ -C ₁₂ -TAD	49	179	167	1855	37.85	10.37	11.12
InPEG ₇₀₀	140	664	288	12 008	86.06	18.07	41.69
InPEG ₇₀₀ -C ₄ -TAD	158	519	308	5772	36.60	11.13	18.75
InPEG ₇₀₀ -C ₆ -TAD	144	567	290	8934	62.14	15.77	30.84
InPEG ₇₀₀ -C ₈ -TAD	63	212	182	4809	76.86	22.69	26.40
InPEG ₇₀₀ -C ₁₂ -TAD	59	230	196	7601	129.47	33.03	38.81



matic C–H and N–H stretching frequencies of the indole unit, supports the incorporation of the indole unit. After modification of the copolymer with hydrophobic TAD derivatives (C_n -TAD), enhanced bands at 2916 cm^{-1} and 2849 cm^{-1} that correspond to the C–H stretching fre-

quency of the alkyl chain (of the TAD unit), along with the appearance of a band at 1702 cm^{-1} , which is attributed to the C=O stretching frequency of the urazole unit formed after the TAD–indole click reaction, support the modification.



Fig. 6 Graphical representation of the structure–antimicrobial activity–hemolytic activity relationship of different PEG derivatives: (A), (C) and (E) represent the effect of different alkyl chain lengths on the antimicrobial activity and selectivity of the InPEG₅₇₅-C_n-TAD polymer series against *P. aeruginosa*, *S. aureus* and *E. coli*, respectively; (B), (D) and (F) represent the effect of PEG chain length on antimicrobial activity and selectivity against *P. aeruginosa*, *S. aureus* and *E. coli*, respectively (the two best-performing polymers InPEG₅₇₅-C₈-TAD and InPEG₇₀₀-C₁₂-TAD were chosen for comparison).



To confirm the cationic nature of the current polymers in solution, their zeta potentials were measured in water (concentration = 0.5 mg mL⁻¹). The zeta potential for the indole-functionalized PEG (InPEG₅₇₅) was found to be +26 mV. After TAD modification (InPEG₅₇₅-C₈-TAD), the zeta potential decreased to +15 mV, confirming the cationic nature of all the polymers. The decrease of zeta potential after TAD modification can be ascribed to the formation of urazole linkages, which are significantly anionic at ambient pH, as their reported pK_a is ~5.3.⁴⁵

The thermal properties of all ten synthesized PEG copolymers were analysed using differential scanning calorimetry (DSC) and thermo-gravimetric analysis (TGA). The TGA thermograms, as depicted in Fig. 4, reveal the thermal stability of the unmodified indole-functionalized PEG polymer (InPEG_x) and hydrophobically modified polymers (InPEG_x-C_n-TAD). A minor weight loss (~5%) was observed around 100 °C (particularly for the more hydrophilic InPEG_x polymers), due to the removal of adsorbed moisture. The thermograms revealed that both InPEG₅₇₅ and InPEG₇₀₀ polymers were stable up to 175 °C (T_{d10}), which is atypical for the PEG backbone. Notably, the hydrophobically modified InPEG_x-C_n-TAD polymers exhibited even greater thermal stability, with a slightly higher decomposition temperature (T_{d10}) of ~200 °C. Furthermore, DSC measurements (Fig. 5 and Fig. S8†) revealed that the glass transition temperature (T_g) values of unmodified InPEG₅₇₅ and InPEG₇₀₀ were -31 °C and -34 °C, respectively. The glass transition temperatures of the TAD derivatives are comparable to those of other similar PEG copolymers that have been reported.⁴⁶ Modifying InPEG_x with a hydrophobic TAD molecule leads to an increase in the T_g values (ranging from 0 to -5 °C) due to hydrogen bonding in the introduced urazole moieties.^{44,47}

2.2. Antimicrobial and hemolytic studies

The antimicrobial activity of all ten polymers was evaluated in solution against three different bacteria – *Pseudomonas aeruginosa* and *Escherichia coli* as model Gram-negative bacteria and *Staphylococcus aureus* as a model Gram-positive bacterium. For quantitative estimation and comparison, the minimum inhibitory concentration (MIC) was determined *via* the broth microdilution method, where the log (CFU mL⁻¹) versus concentration curve was used to determine the MIC. Additionally, a hemolysis test was also performed to determine the HC₅₀, which is the concentration required for 50% RBC lysis. A comparison between the HC and MIC was performed to assess the selectivity of the antimicrobial polymers. The results are reported in Table 2.

For the structure-antimicrobial activity study, two structural variables were examined among the ten polymers (Fig. 6). For two individual sets – InPEG₅₇₅-C_n-TAD and InPEG₇₀₀-C_n-TAD – the PEG spacer length is different, which leads to different periodic structures. For each of these individual sets, 5 different polymers were synthesized with varying alkyl chain lengths (n) – unmodified InPEG_x ($n = 0$), InPEG_x-C₄-TAD ($n = 4$), InPEG_x-C₆-TAD ($n = 6$), InPEG_x-C₈-TAD ($n = 8$), and InPEG_x-C₁₂-TAD ($n = 12$). It is evident that both InPEG₅₇₅ and InPEG₇₀₀ show good antimicrobial activity with excellent selectivity against all three bacteria, far surpassing that of unmodified PEG oligomers (which showed almost no antimicrobial activity). However, InPEG₇₀₀ shows significantly lower hemolytic activity than InPEG₅₇₅, resulting in higher selectivity of InPEG₇₀₀. The higher selectivity can be ascribed to the higher PEG content in its periodic block copolymer structure. The next stage, which involved post-polymer modification of InPEGs, the antimicrobial efficacy initially decreases slightly in the case of C₄-TAD; however, antimicrobial efficacy significantly improved

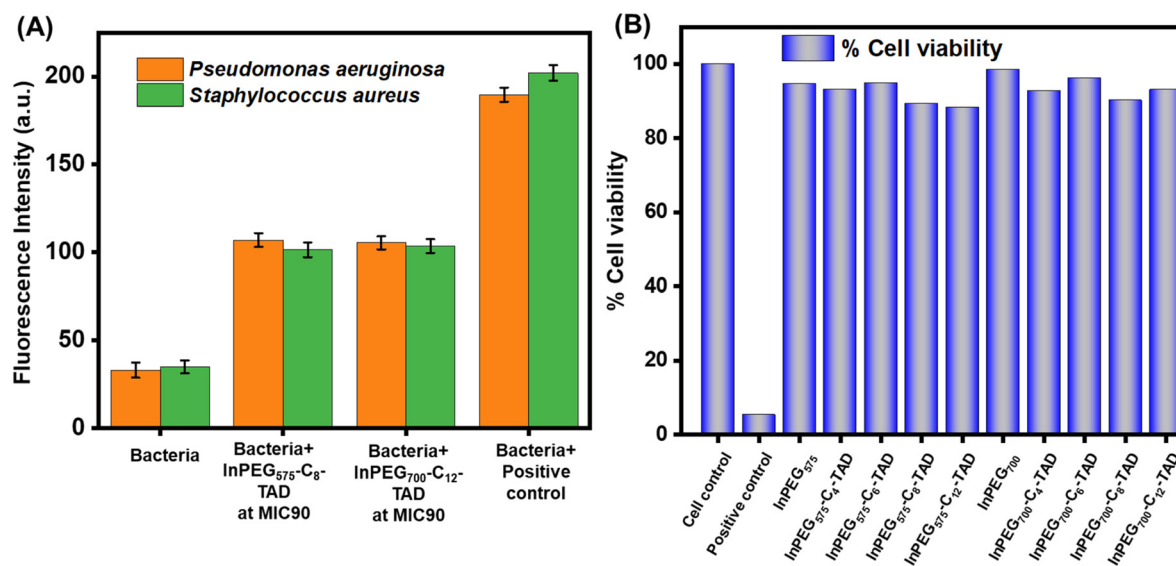


Fig. 7 (A) NPN uptake assay of different bacteria in the presence of InPEG₅₇₅-C₈-TAD and InPEG₇₀₀-C₁₂-TAD. (B) MTT assay assessing the cytotoxicity of different PEG derivatives on HEK293 cells.



when the length of alkyl chains was increased to C-8 and C-12. From the series of 10 polymers, InPEG₇₀₀-C₁₂-TAD was identified as the best antimicrobial agent considering both its MIC and selectivity. The MIC values of InPEG₇₀₀-C₁₂-TAD were 59, 230 and 196 $\mu\text{g mL}^{-1}$ against *P. aeruginosa*, *E. coli* and *S. aureus*, respectively, with excellent selectivity (indicating very low hemolytic activity) of 129, 33 and 39. This further motivated us to calculate the actual hemolytic activity at its MIC. The calculated hemolysis values at 59, 230 and 196 $\mu\text{g mL}^{-1}$ were 3.2, 4.3 and 4.1%, respectively. As all these values are below

5%, they are considered insignificant/null according to the American Society for Testing and Materials (ASTM).^{48,49}

In the next stage, to validate the contact-active/membrane disruption antimicrobial mechanism, we conducted an *N*-phenyl-1-naphthylamine (NPN) uptake assay to determine bacterial membrane permeation by the polymer chains (Fig. 7A). In the literature, the outer membrane assay using 1-*N*-phenyl-1-naphthylamine (NPN) is a well-established method to measure the permeability of bacterial outer membranes.^{50–52} This assay involves incubating bacteria with NPN, where the

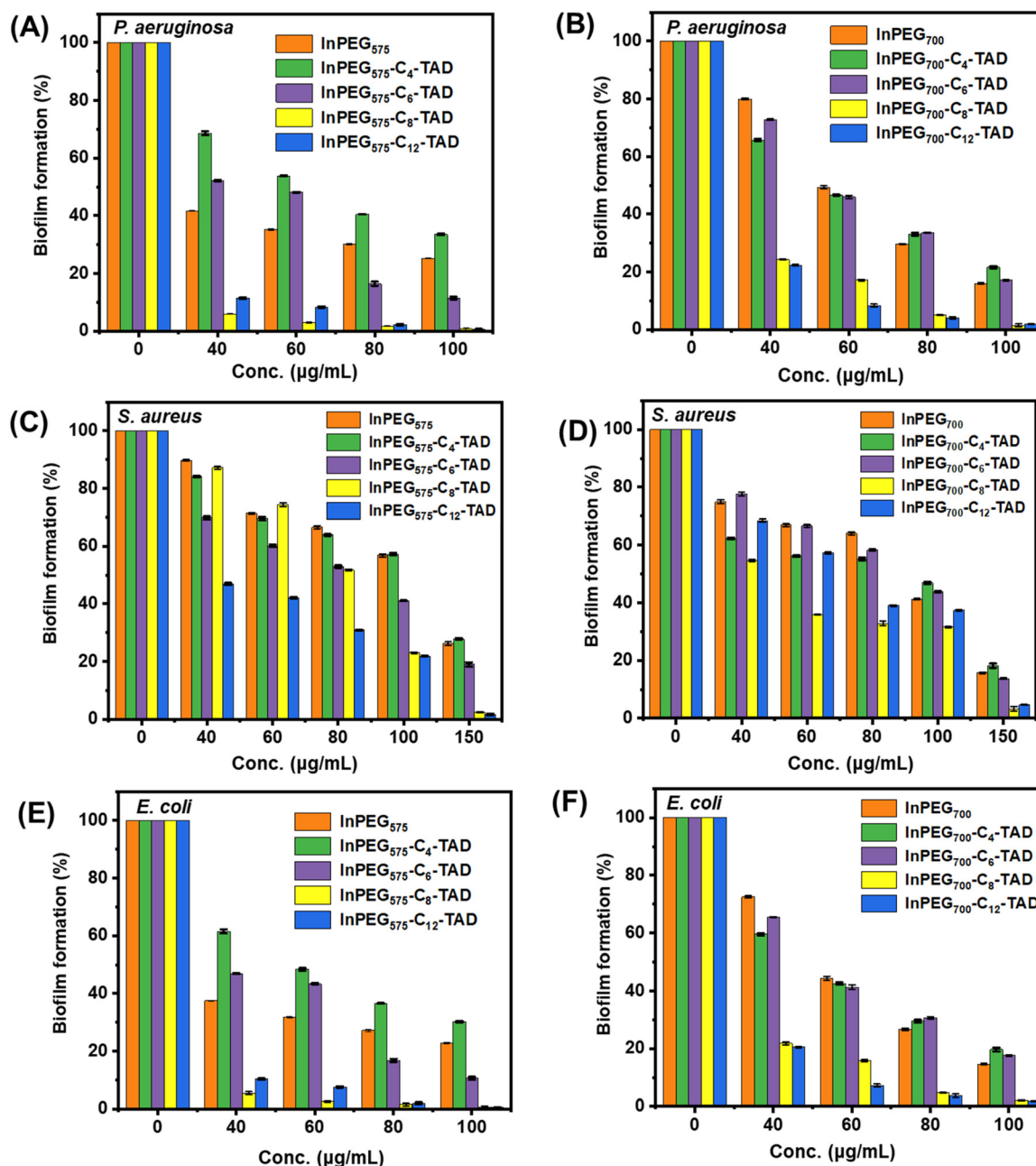


Fig. 8 Crystal violet assay demonstrating the inhibition of biofilm formation in the presence of In-PEG₅₇₅ and its derivatives when tested against *P. aeruginosa*: (A) for the InPEG₅₇₅-C_n-TAD series and (B) for the InPEG₇₀₀-C_n-TAD series; *E. coli*: (C) for the InPEG₅₇₅-C_n-TAD series and (D) for the InPEG₇₀₀-C_n-TAD series; and *S. aureus*: (E) for the InPEG₅₇₅-C_n-TAD series and (F) for the InPEG₇₀₀-C_n-TAD series.



dye binds to the hydrophobic regions of the outer membrane. If membrane integrity is intact, NPN remains outside the cell. However, when the membrane is damaged, NPN enters the cell and exhibits increased fluorescence. Therefore, the intensity of fluorescence correlates with outer membrane permeability, allowing for the assessment of membrane integrity and the effectiveness of antimicrobial agents. The elevated fluorescence intensity following treatment with both InPEG₇₀₀-C₁₂-TAD and InPEG₅₇₅-C₈-TAD demonstrates that the polymers compromise the outer membrane integrity of the bacteria, thereby supporting membrane disruption as the mechanism underlying their antimicrobial effect.

Besides hemolytic activity, cell cytotoxicity is a crucial factor for the application of antimicrobials. To establish the biocompatibility and non-cytotoxic nature of the current polymers, an MTT assay was performed using HEK293 cells (Fig. 7B). Cell cytotoxicity was assessed at a polymer concentration of 1500 $\mu\text{g mL}^{-1}$, which is significantly higher than the reported MIC of the polymers. The results clearly revealed that the polymers are non-toxic, as more than 90% cell viability was noted in all cases.

2.3. Biofilm inhibition assay

Biofilms can be described as communities of bacterial cells enclosed in a hydrated polyanionic exopolysaccharide complex that adheres to a biotic or abiotic surface. Biofilms are particularly important due to their role in persistent infections and antimicrobial tolerance.⁵³ Bacteria can form biofilms on living surfaces such as host tissues and non-living surfaces such as medical devices and implants, and such cells have been proven to be significantly resistant to standard antibiotics and difficult to eliminate.⁵⁴ Therefore, it is important to assess the biofilm inhibition ability of such polymers for their real-life applications.⁵⁵ All ten polymers were assessed using the crystal violet assay (Fig. 8) against three of the most potent biofilm formers, *P. aeruginosa*, *E. coli* (Gram-negative bacteria), and *S. aureus* (Gram-positive bacteria), to determine their ability to inhibit biofilm formation. The results indicate that post-treatment, biofilm inhibition is observed with higher polymer concentrations (please note that in the absence of antimicrobial polymers, biofilm formation reaches 100%, as indicated in Fig. 8, concentration = 0). For most of the polymers, a concentration of 100–150 $\mu\text{g mL}^{-1}$ is able to inhibit biofilm formation (95–100%) in all the bacteria. Although there seems to be no significant difference between the activity of the two individual sets – InPEG₅₇₅-C_n-TAD and InPEG₇₀₀-C_n-TAD – in terms of their structure, the biofilm inhibition activity increases as the length of the alkyl chain increases from C₄ to C₁₂. This trend aligns with the antimicrobial activity patterns observed for these polymers, suggesting that longer alkyl chains enhance both antimicrobial and anti-biofilm properties.

3. Conclusion

In summary, we demonstrated a green and affordable synthetic approach to prepare antimicrobial multifunctional polyethyl-

eneglycols (PEGs) *via* cascade reactions. In the first step, indole-decorated periodic PEG copolymers (InPEG₅₇₅ and InPEG₇₀₀) were prepared *via* aza-Michael-based polyaddition reactions between tryptamine and commercially available PEG diacrylate macromonomers. Here, two different PEG chain lengths influence the polymer periodicity. In the second stage, both the polymers were further post-modified to functionalize them with different alkyl groups *via* TAD–indole reactions, resulting in a series of 10 different multifunctional PEGs. The molecular weight of all the polymers ranged between 5300 and 6600 Da, making them comparable analogues. The glass transition temperature (T_g) of InPEGs was reported to be around $-30\text{ }^\circ\text{C}$, while post-polymerization modifications increased T_g to $(-)$ 4 to $(+)$ 5 $^\circ\text{C}$, mainly due to hydrogen bonding between the incorporated urazole moieties. The antimicrobial activity of all the polymers was evaluated against three different bacteria, *P. aeruginosa*, *E. coli* and *S. aureus*. Antimicrobial and hemolytic studies revealed that structure plays a pivotal role in tuning the antimicrobial efficacy and selectivity (HC/MIC) of the polymers. The selectivity (HC/MIC) reported for the best prototype (InPEG₇₀₀-C₁₂-TAD) was 129, 33 and 39 against *P. aeruginosa*, *E. coli* and *S. aureus*, respectively. The NPN assay revealed bacterial membrane permeation by the polymers as the major bactericidal mechanism. All the polymers were non-cytotoxic, as revealed by the MTT assay. Furthermore, biofilm inhibition analysis of the polymers using the crystal violet assay revealed that both C₈- and C₁₂-functionalized analogues exhibited significant biofilm inhibition (>90%) above 60 $\mu\text{g mL}^{-1}$, whereas complete inhibition was noted (\sim 99%) at 100 $\mu\text{g mL}^{-1}$ for *P. aeruginosa* and *E. coli* and at 150 $\mu\text{g mL}^{-1}$ for *S. aureus*.

Author contributions

SK: synthesis, characterization, relevant data analysis (lead), and writing (lead); AH: biological experiments and analysis (lead) and writing (supporting); AA: synthesis, characterization (supporting), and writing (supporting); OM: supervision (biological experiments) and writing (supporting and review); and SC: conceptualization of the actual work and supervision (synthesis and relevant characterization) and writing (lead and review). All the authors have approved the final manuscript.

Data availability

The data supporting this article have been included as part of the ESI.†

Conflicts of interest

There are no conflicts to declare.



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References

- R. Laxminarayan, A. Duse, C. Wattal, A. K. M. Zaidi, H. F. L. Wertheim, N. Sumpradit, E. Vlieghe, G. L. Hara, I. M. Gould, H. Goossens, C. Greko, A. D. So, M. Bigdeli, G. Tomson, W. Woodhouse, E. Ombaka, A. Q. Peralta, F. N. Qamar, F. Mir, S. Kariuki, Z. A. Bhutta, A. Coates, R. Bergstrom, G. D. Wright, E. D. Brown and O. Cars, *Lancet Infect. Dis.*, 2013, **13**, 1057–1098.
- M. A. Kohanski, D. J. Dwyer and J. J. Collins, *Nat. Rev. Microbiol.*, 2010, **8**, 423–435.
- K. U. Jansen, C. Knirsch and A. S. Anderson, *Nat. Med.*, 2018, **24**, 10–19.
- A. Fleming, *Br. Med. Bull.*, 1944, **2**, 4–5.
- A. Tomasz, *Annu. Rev. Microbiol.*, 1979, **33**, 113–137.
- C. D. Fjell, J. A. Hiss, R. E. W. Hancock and G. Schneider, *Nat. Rev. Drug Discovery*, 2012, **11**, 37–51.
- K. A. Brogden, *Nat. Rev. Microbiol.*, 2005, **3**, 238–250.
- M. Zasloff, *Nature*, 2002, **415**, 389–395.
- M. N. Melo, R. Ferre and M. A. R. B. Castanho, *Nat. Rev. Microbiol.*, 2009, **7**, 245–250.
- H. Leontiadou, A. E. Mark and S. J. Marrink, *J. Am. Chem. Soc.*, 2006, **128**, 12156–12161.
- R. E. W. Hancock and H.-G. Sahl, *Nat. Biotechnol.*, 2006, **24**, 1551–1557.
- R. Spohn, L. Daruka, V. Lázár, A. Martins, F. Vidovics, G. Grézal, O. Méhi, B. Kintsés, M. Számel, P. K. Jangir, B. Csörgő, Á. Györkei, Z. Bódi, A. Faragó, L. Bodai, I. Földesi, D. Kata, G. Maróti, B. Pap, R. Wirth, B. Papp and C. Pál, *Nat. Commun.*, 2019, **10**, 4538.
- E. A. Porter, B. Weisblum and S. H. Gellman, *J. Am. Chem. Soc.*, 2002, **124**, 7324–7330.
- Y. Wu, G. Xia, W. Zhang, K. Chen, Y. Bi, S. Liu, W. Zhang and R. Liu, *J. Mater. Chem. B*, 2020, **8**, 9173–9196.
- E.-R. Kenawy, S. D. Worley and R. Broughton, *Biomacromolecules*, 2007, **8**, 1359–1384.
- M. Haktaniyan and M. Bradley, *Chem. Soc. Rev.*, 2022, **51**, 8584–8611.
- Y. Ko, V. K. Truong, S. Y. Woo, M. D. Dickey, L. Hsiao and J. Genzer, *Biomacromolecules*, 2022, **23**, 424–430.
- A. Khlyustova, M. Kirsch, X. Ma, Y. Cheng and R. Yang, *J. Mater. Chem. B*, 2022, **10**, 2728–2739.
- P. R. Judzewitsch, T.-K. Nguyen, S. Shanmugam, E. H. H. Wong and C. Boyer, *Angew. Chem., Int. Ed.*, 2018, **57**, 4559–4564.
- P. Pham, S. Oliver, E. H. H. Wong and C. Boyer, *Polym. Chem.*, 2021, **12**, 5689–5703.
- P. T. Phuong, S. Oliver, J. He, E. H. H. Wong, R. T. Mathers and C. Boyer, *Biomacromolecules*, 2020, **21**, 5241–5255.
- A. O. Biying, M. J. H. Ong, W. Zhao, N. Li, H.-K. Luo and J. M. W. Chan, *ACS Appl. Polym. Mater.*, 2024, **6**, 8169–8177.
- S. Chattopadhyay, E. Heine, A. Mourran, W. Richtering, H. Keul and M. Möller, *Polym. Chem.*, 2016, **7**, 364–369.
- S. Chattopadhyay, E. T. Heine, H. Keul and M. Möller, *Macromol. Biosci.*, 2014, **14**, 1116–1124.
- S. Chattopadhyay, H. Keul and M. Moeller, *Green Chem.*, 2013, **15**, 3135–3139.
- J. Zhang, Z. Yang, Y.-H. Li, S. Durrani, A.-P. Pang, Y. Gao, F.-G. Wu and F. Lin, *ACS Appl. Polym. Mater.*, 2022, **4**, 425–434.
- P. M. Alves, R. F. Pereira, B. Costa, N. Tassi, C. Teixeira, V. Leiro, C. Monteiro, P. Gomes, F. Costa and M. C. L. Martins, *ACS Appl. Polym. Mater.*, 2022, **4**, 5012–5026.
- M. Másson, *Carbohydr. Polym.*, 2024, **337**, 122159.
- B.-I. Andreica, L. Mititelu-Tartau, I. Rosca, I. M. Pelin, E. Nicol and L. Marin, *Carbohydr. Polym.*, 2024, **342**, 122389.
- S. Chattopadhyay, E. Heine, H. Keul and M. Moeller, *Polymers*, 2014, **6**, 1618–1630.
- C. Krumm, M. Hijazi, S. Trump, S. Saal, L. Richter, G. G. F. K. Noschmann, T.-D. Nguyen, K. Preslikoska, T. Moll and J. C. Tiller, *Polymer*, 2017, **118**, 107–115.
- M. Concilio, R. Garcia Maset, L. P. Lemonche, V. Kontrimas, J.-I. Song, S. K. Rajendrakumar, F. Harrison, C. R. Becer and S. Perrier, *Adv. Healthcare Mater.*, 2023, **12**, 2301961.
- Y. Hu, J. Zhao, M. Yang, X. Wang, H. Zhang, J. Zhang, Z. Zhu and J. Rao, *ACS Appl. Polym. Mater.*, 2022, **4**, 4868–4875.
- E. N. Şimşek, A. Akdağ and P. Z. Çulfaz-Emecen, *Polymer*, 2016, **106**, 91–99.
- M. Kim, W. Mun, W. H. Jung, J. Lee, G. Cho, J. Kwon, D. J. Ahn, R. J. Mitchell and B.-S. Kim, *ACS Nano*, 2021, **15**, 9143–9153.
- J. D. Katiyar, A. Halder, M. Avais, H. Aidasani, O. Mukherjee and S. Chattopadhyay, *ACS Appl. Polym. Mater.*, 2023, **5**, 9742–9750.
- X. Li, S. İlk, Y. Liu, D. B. Raina, D. Demircan and B. Zhang, *Polym. Chem.*, 2022, **13**, 2307–2319.
- X. Li, X. Wang, S. Subramanian, Y. Liu, J. Rao and B. Zhang, *Biomacromolecules*, 2021, **23**, 150–162.
- S. Billiet, K. De Bruycker, F. Driessen, H. Goossens, V. Van Speybroeck, J. M. Winne and F. E. Du Prez, *Nat. Chem.*, 2014, **6**, 815–821.
- S. Kumari, M. Avais and S. Chattopadhyay, *Polymer*, 2022, **256**, 125219.
- S. Kumari and S. Chattopadhyay, *Surf. Interfaces*, 2024, **46**, 103884.
- A. Anand, R. K. Bhagat, S. Ghosh and S. Chattopadhyay, *ACS Appl. Polym. Mater.*, 2024, **6**, 11487–11496.
- L. Vlaminck, B. Van de Voorde and F. E. Du Prez, *Green Chem.*, 2017, **19**, 5659–5664.
- S. Kumari and S. Chattopadhyay, *Mol. Syst. Des. Eng.*, 2024, **9**, 490–499.



- 45 S. Chattopadhyay and F. Du Prez, *Eur. Polym. J.*, 2016, **81**, 77–85.
- 46 M. Avais and S. Chattopadhyay, *Macromol. Chem. Phys.*, 2022, **223**, 2200109.
- 47 L. Vlamincx, K. De Bruycker, O. Türünç and F. E. Du Prez, *Polym. Chem.*, 2016, **7**, 5655–5663.
- 48 U. Bernauer, L. Bodin, Q. Chaudhry, P. J. Coenraads, M. Dusinska, E. Gaffet, E. Panteri, V. Rogiers, C. Rousselle, M. Stepanik, T. Vanhaecke, S. Wijnhoven, N. von Goetz, W. H. de Jong and A. Simonnard, *Regul. Toxicol. Pharmacol.*, 2020, **112**, 104611.
- 49 R. Luna-Vázquez-Gómez, M. E. Arellano-García, J. C. García-Ramos, P. Radilla-Chávez, D. S. Salas-Vargas, F. Casillas-Figueroa, B. Ruiz-Ruiz, N. Bogdanchikova and A. Pestryakov, *Materials*, 2021, **14**, 2792.
- 50 F. Chen, Y. Tang, H. Zheng, Y. Xu, J. Wang and C. Wang, *ChemMedChem*, 2019, **14**, 1457–1465.
- 51 J. Wang, S. Chou, L. Xu, X. Zhu, N. Dong, A. Shan and Z. Chen, *Sci. Rep.*, 2015, **5**, 15963.
- 52 I. M. Helander and T. Mattila-Sandholm, *J. Appl. Microbiol.*, 2000, **88**, 213–219.
- 53 T.-F. C. Mah and G. A. O'Toole, *Trends Microbiol.*, 2001, **9**, 34–39.
- 54 P. Keshvardoust, V. A. A. Huron, M. Clemson, F. Constancias, N. Barraud and S. A. Rice, *npj Biofilms Microbiomes*, 2019, **5**, 22.
- 55 Q. Li, L. Wang, L. Yu, C. Li, X. Xie, H. Yan, W. Zhou, C. Wang, Z. Liu, G. Hou and Y.-Q. Zhao, *Biomacromolecules*, 2024, **25**, 1180–1190.

