#### **REVIEW**

# **Novel Strategies to Combat Bacterial Bioflms**

Fatemeh Hemmati<sup>1,2</sup> · Mohammad Ahangarzadeh Rezaee<sup>2</sup> · Saba Ebrahimzadeh<sup>3</sup> · Leila Yousefi<sup>1,4</sup> · **Roghayeh Nouri1,2 · Hossein Samadi Kafl4 · Pourya Gholizadeh[4](http://orcid.org/0000-0002-1451-3996)**

Received: 9 January 2021 / Accepted: 9 April 2021

© The Author(s), under exclusive licence to Springer Science+Business Media, LLC, part of Springer Nature 2021

## **Abstract**



Bioflms are considered as a severe problem in the treatment of bacterial infections; their development causes some noticeable resistance to antibacterial agents. Bioflms are responsible for at least two-thirds of all infections, displaying promoted resistance to classical antibiotic treatments. Therefore, fnding new alternative therapeutic approaches is essential for the treatment and inhibition of bioflm-related infections. Therefore, this review aims to describe the potential therapeutic strategies that can inhibit bacterial bioflm development; these include the usage of antiadhesion agents, AMPs, bacteriophages, QSIs, aptamers, NPs and PNAs, which can prevent or eradicate the formation of bioflms. These antibioflm agents represent a promising therapeutic target in the treatment of bioflm infections and development of a strong capability to interfere with diferent phases of the bioflm development, including adherence, polysaccharide intercellular adhesion (PIA), quorum sensing molecules and cell-to-cell connection, bacterial aggregation, planktonic bacteria killing and host-immune response modulation. In addition, these components, in combination with antibiotics, can lead to the development of some kind of powerful combined therapy against bacterial bioflm-related infections.

**Keywords** Bioflm · Antibioflm agents · Bioflm-related infections · Therapeutic approaches

# **Introduction**

Most nosocomial infections have been caused by opportunistic pathogens in the recent decades; these are associated with various infections including bacteremia, urinary tract infections (UTIs), wound, meningitis and bioflm-associated infections [[1\]](#page-11-0). Bacteria that are frmly attached to artifcial medical devices cause bioflm-associated infections. Treatment of these infections is difficult because of the increased antibiotic resistance in the bioflm, which is commonly due to multidrug-resistant strains [[2,](#page-11-1) [3](#page-11-2)]. Nowadays, bioflm

 $\boxtimes$  Pourya Gholizadeh poorya.gholizadeh@gmail.com

- Student Research Committee, Tabriz University of Medical Sciences, Tabriz, Iran
- <sup>2</sup> Immunology Research Center, Tabriz University of Medical Sciences, Tabriz, Iran
- <sup>3</sup> Department of Food Science and Technology, Faculty of Agriculture and Natural Resources, Urmia University, Urmia, Iran
- <sup>4</sup> Drug Applied Research Center, Tabriz University of Medical Sciences, Tabriz, Iran

infections have become a serious concern and a main global healthcare problem [[1\]](#page-11-0). The identification of novel therapeutic targets to fght bioflm-related infections, thus, signifes one of the primary problems in the feld of antibiotic therapy [[4\]](#page-11-3). Bacteria are present in the environment in two forms: free-living or bioflm. Bacteria inside a bioflm can demonstrate low levels of sensitivity to some antimicrobial compounds including biocides and antibiotics [[2](#page-11-1), [3\]](#page-11-2). Therefore, antibioflm compounds can be regarded as one of the most encouraging options to combat bioflm-related infections. These compounds have shown a broad range of biological functions including antibacterial properties on free-living cells, as well as antibioflm properties [\[5](#page-11-4)].

Bioflm is known as a bacterial population in which cells adhere to biotic or abiotic surfaces by extracellular polymeric substances (EPS). Bioflm could be considered as a strategy some bacteria could adopt to withstand adverse conditions such as desiccation, host defense system and antibacterial agents [\[6](#page-11-5)].

The stages involved in the formation of bioflm in bacteria comply with a generic model consisting of three stages. The features of each stage are characteristics in diferent species, but the general traits are all alike [[7\]](#page-11-6). The main

stages in the bioflm formation are attachment, accumulation and dispersal [[8](#page-11-7)]. In the frst stage, the attachment to the surface is reversible and not very strong. Next, a loose bond is established between cells, and the extracellular matrix begins to secrete. In the later stages, the bioflm structure gains shape and other bacteria still in the environment attach themselves to this structure. Following the evolution of the bioflm structure, the cells are released back into the environment, causing new bioflm foci elsewhere [[9](#page-11-8)].

In the bioflm phase, bacteria are surrounded in a selfproduced extracellular matrix principally made of EPS [[10](#page-11-9)]. EPS comprises over 90% of the dry weight of the bioflm, facilitating bonding to surfaces, which can cause microcolony formation and resistance to antimicrobial agents [[11\]](#page-11-10). EPS is made up of exopolysaccharides, proteins and extracellular DNA (eDNA) [[10\]](#page-11-9). EPS matrix reduces the permeation of antibiotics to spread bacteria within the bioflm by dispersion restriction or neutralization of antibacterial agents with extracellular polysaccharides [[12\]](#page-11-11). Bacterial bioflms are signifcant in causing infectious diseases, especially chronic infections, in the host [[13,](#page-11-12) [14\]](#page-11-13). According to some estimations previously carried out, approximately 80% of bacterial infections in the human body are associated with the formation of bioflms, which can increase the mortality rate in the hospitalized patients [[2](#page-11-1), [14\]](#page-11-13). Therefore, managing bioflm-related infections is challenging because of the problems inherent in inhibiting and treating them [\[3](#page-11-2)].

Given the specifc conditions in the bioflm environment, treating and employing antibiotics to tackle this problem can be regarded as one of the important challenges the medical sciences face. Bioflm-associated infections cannot be treated by classical antibiotics, and they are a challenge worldwide. This has led to a set of studies aiming to fnd modern treatments for biofilm infections [[15\]](#page-11-14). Therefore, new antibacterial or antibiofilm strategies with various mechanisms of action are urgently needed. Among these strategies, development of new classes of antimicrobial peptides (AMPs) [\[16](#page-11-15)], bacteriophages [[17\]](#page-11-16), quorum sensing inhibitors (QSIs) [[18\]](#page-11-17), aptamers [[19\]](#page-11-18), nanoparticles (NPs) [\[20\]](#page-11-19) and peptide nucleic acids (PNAs) [[21\]](#page-11-20) can be considered as the most feasible solution.

In the recent years, several innovative antibioflm agents have been developed to limit bacterial adhesion to abiotic and biotic surfaces; these are intended to target bacterial signals for removing grown bioflms or displacing cells from the established bioflms [\[4](#page-11-3)]. Therefore, this review aims to review and describe the potential therapeutic strategies that could be applied to prevent bacterial bioflm development; these include the usage of antiadhesion agents, AMPs, bacteriophages, QSIs, aptamers, NPs and PNAs, which can prevent or eradicate the bioflm formation.

# **Properties of Desirable Antibioflm Agents**

Bioflms are bacterial populations demonstrating exclusive properties in comparison with their free-living forms [[22\]](#page-11-21). These properties should be correctly considered when assessing the potential of bioflm inhibition mechanisms. The EPS matrix in the bioflm development plays a signifcant and critical role in determining antibiotic resistance mechanisms of bioflm [[4\]](#page-11-3). It basically establishes a dispersion barrier which can prevent the interaction of antibacterial agents with bacterial cells [[23\]](#page-11-22). Restricting or preventing EPS accumulation and having the capability to permeate EPS could be regarded as characteristics of an ideal antibioflm agent [[24](#page-11-23)]. Therefore, characteristics of an ideal antibioflm substance include antibacterial activities, easy penetration into the cell, unique structure, interference with the machinery of bacterial cells communication and synergism with other antibacterial agents [\[4](#page-11-3)]. Many of such properties can be observed in natural and synthetic antibioflm substances  $[25-28]$  $[25-28]$  $[25-28]$ . Inhibition of the biofilm development by these compounds may simplify the treatment of bioflmrelated infections.

# **Strategies for the Inhibition and Disruption of Bacterial Bioflms**

Bioflms are highly resistant to conventional antibiotics; therefore, new alternative therapeutic approaches are needed to treat bioflm-related infections [[3,](#page-11-2) [29](#page-12-1)]. There are several strategies to inhibit and eradicate bioflm development; these include antiadhesion agents, AMPs, bacteriophages, QSIs, aptamers, NPs and PNAs.

### **Antiadhesion Agents**

The introduction of bacteria to a surface is infuenced by the stochastic process, which is driven by gravitational forces, the surrounding hydrodynamic forces and Brownian motion [[30,](#page-12-2) [31](#page-12-3)]. To overcome repulsive and hydrodynamic forces, motile bacteria utilize fagella, which can play an important role in the initial attachment in the case of several pathogens such as *Escherichia coli*, *Pseudomonas aeruginosa*, *Listeria monocytogenes* and *Vibrio cholera* [\[32–](#page-12-4)[35](#page-12-5)]. The roles of chemotaxis in directing attachment and bioflm formation in response to nutrient composition have been demonstrated. Schmidt et al. [[36\]](#page-12-6), for instance, showed that the CheR1 methyltransferase mutations of *P. aeruginosa* could alter the amino acid response and inhibit bioflm maturation through impairing attachment. In addition, bioflm defects have been revealed in the *tar* gene (encodes methyl-accepting chemotaxis protein II) in uropathogenic *E. coli* (UPEC) [\[37\]](#page-12-7). Type 1 pili is another type of adhesion in UPEC and other *E. coli* that can play a role in the initial attachment, maintaining a steadfast grip on the surface and shear forces [[38,](#page-12-8) [39](#page-12-9)]. These pili, which are assembled by the chaperon usher pathway (CUP), are multi-subunit adhesive [[40\]](#page-12-10). CUP pili systems facilitate adherence in a niche-specifc manner [[37,](#page-12-7) [40,](#page-12-10) [41](#page-12-11)]. Adherence in the type 1 pili is mediated by the FimH adhesion, which recognizes mannose-rich regions [\[42](#page-12-12)]. Antigen 43 and curli fbers are other types of adhesion mediating interbacterial interactions and attachment on biotic and abiotic surfaces [[43](#page-12-13)]. There are other attachment organelles in bacteria, including type IV pili and numerous CUP fmbriae (such as CupA) in *P. aeruginosa* [[44\]](#page-12-14), Ace, Esp and Ebp in *Enterococcus faecalis* [[45\]](#page-12-15), and SagA and Acm in *Enterococcus faecium* [\[46](#page-12-16)]. Therefore, several studies have focused on the development of compounds interacting and interfering with the frst step of attachment, leading to the bioflm formation. Mannosides are molecules that competitively inhibit FimH mannose binding [\[47\]](#page-12-17). Among mannosylated proteins, the highest afnity of the monovalent ligands has been displayed in long chain aryl mannosides and alkyl mannosides, which could be due to the increased interactions with Tyr-48, Tyr-137 and Ile-52 of the binding pocket [[48\]](#page-12-18). The development of monomeric biphenyl mannosides could lead to FimH inhibitors [\[49\]](#page-12-19), which can prevent in vitro UPEC bioflm formation and interfere with the in vivo adherence and invasion of UPEC [\[49](#page-12-19)]. Cusumano et al. [[49\]](#page-12-19) also demonstrated that

<span id="page-2-0"></span>**Table 1** Peptides with the antibioflm activity

the combination of mannosides with TMP-SMZ (trimethoprim-sulfamethoxazole) could enhance antimicrobial treatment in clinical settings. In addition, the bladder bacterial load of UPEC was reduced in orally mannoside treated mice within 6 h, thus serving as an efficient therapeutic strategy for chronic UTIs [\[49](#page-12-19)].

In parallel, pilicides have been introduced to inhibit CUPs and type 1 pili, which are components derived from peptodomimetic scafolds such as C-2 substituted thiazolo and dihydrothiazolo ring-fused 2-pyridones, as well as the bromomethyl substituted scafold that can interfere and interact with the exportation of the subunits corresponding to the pili structure [[50](#page-12-20), [51\]](#page-12-21). Furthermore, curlicides (components derived from ring-fused 2-pyridones, such as FN075 and BibC6) have been introduced to inhibit the curli synthesis followed by the inhibition of the bioflm formation [\[43](#page-12-13)]. According to some studies, bacteria utilize a variety of receptors and adhesion types during their adherence process, facilitating adhesion in a niche-specifc manner; therefore, interactions of multiple molecules may need to be inhibited to remove a bioflm from surfaces.

# **Antimicrobial Peptides**

Antimicrobial peptides (AMPs) are small cationic molecules displaying a broad range of activity against microorganisms [[52\]](#page-12-22). AMPs are divided into natural and synthetic groups. Table [1](#page-2-0) shows the list of AMPs and their properties. Natural AMPs are produced by cellular tissues in a wide range of



organisms; they can be a source for synthetic AMPs [\[53,](#page-12-23) [54](#page-12-24)]. AMPs are suggested as a potential method to cope with the bioflm formation. AMPs as antibioflm agents have received much attention. Many AMPs have a positive charge, enabling them to interact with the phosphate groups of lipopolysaccharides as negatively charged components of the cell membrane in Gram-negative bacteria or lipoteichoic acids in Gram-positive bacteria [\[55](#page-12-25), [56\]](#page-12-26). Most AMPs exhibit strong antibioflm properties against antibiotic-resistant bacteria, which can be efective with diferent mechanisms at various phases of the bioflm development and on diferent molecular targets [\[16\]](#page-11-15). Some peptides can kill bacteria through membrane distraction and/or pore formation, as well as inhibition of the bacterial cell division; further, they can be effective on bacteria through preventing the adherence of the bacterial cells to the substrate surface, downregulating QS signals and removing the pre-formed bioflm [[16](#page-11-15)]. Downregulation of genes involved in the motility and inhibition of a series of cellular biological procedures, such as the synthesis of cell walls, DNA, RNA and proteins, are other instances of AMP-antibioflm mechanisms of function [\[57](#page-12-27)[–60](#page-13-10)]. Figure [1](#page-3-0) shows multiple mechanisms of antimicrobial peptides on the bacterial bioflm formation.

Antibioflm peptides (ABPs) are a type of AMPs with the antibioflm activity; they prevent the bioflm development at concentrations much lower than those of common antibiotics. The minimum inhibitory concentration (MIC) is, in fact, greater than the minimum bioflm inhibitory concentration (MBIC) [\[71](#page-13-11)]. ABPs can be linked to ppGpp molecules (second messenger nucleotide), disrupting them; they are involved in several metabolic pathways like colonization, attachment and aggression in bacteria. Therefore, ABPs prevent the increase of these molecules in the cell and inhibit the bioflm development in bacteria [\[72,](#page-13-12) [73](#page-13-13)]. ABPs commonly interact with signal molecules to apply control or distraction efects on the bioflms development; therefore, they can help other antibacterial agents to cope with the bacterial cells. For instance, Ribeiro et al. [[74\]](#page-13-14) described that ABPs promoted the sensitivity of carbapenemase-producing *Klebsiella pneumoniae* to carbapenems.

It has been demonstrated that most AMPs can be in combined with antibiotics to enhance their ability to prevent bioflm development and eliminate mature bioflms. Synergism of AMPs with other antibacterial agents is promised in the cleaning of bioflm-related infections, as they can increase the antibiofilm effect and reduce the drug dose [\[68](#page-13-7)]. Tong et al. [[75](#page-13-15)] also reported the synergistic interaction between nisin (a natural AMP) and penicillin or chloramphenicol against the *E. faecalis* biofilm. The synergistic efficacy of nisin with nafcillin on bioflm in *Streptococcus mutans* was evaluated by Tong et al. [[76](#page-13-16)]. Their results showed that nisin in combination with antibiotics considerably decreased the bioflm development. Therefore, AMPs could be regarded as potential antibioflm agents through diferent mechanisms



<span id="page-3-0"></span>**Fig. 1** Multiple mechanisms of antimicrobial peptides on the bacterial bioflm formation

including functional inhibition of proteins, cell-membranedisrupting action, detoxifcation of lipoteichoic acid and lipopolysaccharide, and binding with DNA.

#### **Bacteriophages**

The viruses that can infect and kill bacteria are known as bacteriophages (phages); they are not able to infect human or animal cells [[77\]](#page-13-17). Therefore, phages are known as the key hunters of bacteria in the environment, and phage therapy has always been of interest for medical goals. Phage therapy emphasizes on lytic phages because they kill their host bacterial cells. In addition, lytic phages lack integrases and/or genes that participate in the horizontal gene transfer. Lytic phages are capable of lysing their host bacterial cells; as well, they can be amplifed inside the host bacterial cells. Lysis of the host bacterial cells destroys the bacteria, as well as releasing the progeny phages for the re-infection of more bacteria. Phages are species-specifc; thus, they can be used to target special pathogenic bacteria, without any efect on the commensal bacteria [\[17](#page-11-16)].

Phage therapy has several advantages in comparison to common antibiotics; these include specifcity of function, a narrow range of action, greater safety, greater tolerability, an efect limited to the site of infection and cost efectiveness [[78](#page-13-18)]. The use of phages to inhibit bacterial infections is being followed as an alternative to antibiotics [\[79](#page-13-19)]. According to some recent studies, the role of phages in the eradication and/or prevention of bacteria bioflms has been highlighted. Phages are capable of penetrating into the struc-ture of bacterial biofilms and removing them [\[80](#page-13-20), [81\]](#page-13-21). There are several water channels in the bioflm structure, allowing the phages to easily penetrate into the bioflms' inner structure [\[82\]](#page-13-22). Furthermore, most of the phages generate depolymerases, which are capable of hydrolysing the EPS of the bacterial bioflms [\[83\]](#page-13-23). For instance, phage phi 15 produces the polysaccharide depolymerase enzyme, which hydrolyses the EPS of *Pseudomonas putida* and inhibits its bioflm formation [[84](#page-13-24)]. As well, it has been observed that T4-like and Φ29-like phages from the family of *Myoviridae* and *Podoviridae*, respectively, suppress the bioflm of *Staphylococcus aureus* [\[85](#page-13-25)].

Phage-derived enzymes like lysin enzymes are known as bactericidal components that hydrolase peptidoglycan, the key composition of the cell-wall of both Gram-negative and Gram-positive bacteria. Destruction of peptidoglycan by lysine could induce lysis and degrade the bacterial cell-wall and bioflm structure [\[86](#page-13-26), [87\]](#page-13-27). Current studies have shown that phage-derived lysin enzymes can potentially act as antibioflm development and antibacterial components [\[81,](#page-13-21) [88,](#page-13-28) [89\]](#page-13-29). It has also been reported that osmotic lysis, independent of bacterial metabolism, can occur upon applying Art-175 lysin against *P. aeruginosa* bioflms. This phenomenon is often important for the elimination of the bacterial bioflms; owing to lysin, even at low metabolic levels, persistent bacteria inside bioflms can be killed [\[90](#page-13-30)].

Some phages cannot produce specifc enzymes to permeate and difuse into the EPS matrix for the bioflms inhibition [[91\]](#page-13-31). Nevertheless, phages are often genetically modifed to produce enzymes that can damage the EPS matrix and ease the elimination of bioflms. Lu et al. [[80\]](#page-13-20) demonstrated that a modifed phage T7 of *E. coli* intracellularly produced a hydrolase during infection, which could increase the bioflm removal by being released into the extracellular matrix. An eradication rate more than 99% was observed by testing on *E. coli* bioflms, thus confrming the advantage of using the modifed phages.

In phage therapy, individual or cocktail (mixture) phages can be used. The usage of cocktails versus individual phages could considerably increase the host spectrum and decrease the generation of phage-resistant types. The cocktails of phages have been confrmed to be efective in the prevention of the bioflm development and bioflm removal. Antibacterial components such as disinfectants and antibiotics may be applied with phages to increase the efficiency of the biofilm eradication [[81](#page-13-21)].

The combination therapy of phages and antibiotics can increase the treatment efectiveness, as well as inhibiting the resistance to phages without enhancing the toxicity of the antibiotics [[79\]](#page-13-19). A combined usage of phages with antibiotics increases the synergistic activation of these factors in improving the bioflm disruption. Combined therapy of phage T4 and tobramycin on *E. coli* bioflms has also been done; this could drastically decrease tobramycin-resistant *E. coli*. The same experiment was reported against *P. aeruginosa* bioflms by phage PB-1 [[92\]](#page-13-32). Also, the synergistic efficacy of the phage with amoxicillin on the biofilm of *Klebsiella pneumoniae* B5055 was demonstrated by Bedi et al. [[79](#page-13-19)]. The combined therapy of amoxicillin and phages signifcantly increased the elimination of the bioflm development in the case *K. pneumoniae*, as compared to each of the agents alone [[79](#page-13-19)]. Henriksen et al. [\[93](#page-14-0)] also demonstrated that sub-MIC concentrations combined ciprofoxacin and phages have a synergic efect on *P. aeruginosa* bioflm, showing a decrease of about 6 log in the bioflm formation. Depolymerized enzymes in combination with antibiotics could promote the antibacterial efficacy by facilitating the entrance of antibiotics to the bioflm inside. These enzymes could diminish the adherence of EPS matrix and bacteria, thus favoring the function of antibiotics [\[94](#page-14-1)].

However, a main limitation of the combined therapy of phages and antibiotics is that antibiotic-resistant bioflms may be increased because phages could favorably infect antibiotic-sensitive bacteria [[17\]](#page-11-16). Furthermore, the interference with bacterial metabolism is needed for the DNA replication and protein synthesis of phages. Since phage therapy

is strictly associated to the growth condition of its bacterial host and common antibiotics potentially act on the log phase of bacterial growth, which have the maximum metabolism, it can be regarded as one of the obstacles in combining antibiotics and phage therapy [\[81](#page-13-21), [95,](#page-14-2) [96\]](#page-14-3). Therefore, to prevent incompatibility, the potential adverse effects of the combination therapy of antibiotics should be considered. As some pathogenic bacteria prefer to remove competitors, the use of the combination therapy of antibiotics and phage cocktails is especially interesting for the treatment of mixed infections.

#### **Quorum Sensing Inhibitors**

Bacterial cell-to-cell communication is known as Quorum sensing (QS); it directly acts in the bioflm development of various bacterial species. This system can control the expression of various pathogenic and virulence genes in the bioflm phase [[97,](#page-14-4) [98\]](#page-14-5). In this system, small molecules, known as autoinducers, are responsible for the communication of bacteria with each other. The gene expression levels may display major changes when the bacterial density reaches the concentration threshold of autoinducers [[99](#page-14-6)]. Changes in the gene expression levels could affect (induction or suppression) various virulence factors in bacteria; these also include the bioflm production. The changes in the environment surrounding the microorganism can modify the planktonic state to become a bioflm. The gene expression in the planktonic state undergoes many changes during the transition to the bioflm state. Cell surface molecules, specifc metabolic pathways, and the production of various factors can all contribute to the bacterial survival under bioflm conditions [[44\]](#page-12-14). Quorum quenchings (QQ) or quorum sensing inhibitors (QSIs) are described as molecules produced by eukaryotes and/or prokaryotes with the ability to prevent the QS systems, which can lead to decreasing the expression of efflux pump genes and the disruption of bacterial bioflms [[99\]](#page-14-6). Several methods have been applied to disturb QS, such as blocking the production of acyl-homoserine lactones (AHLs), diminishing the activity of the AHL synthase, disturbing and inactivating AHLs, and utilizing numerous competitors compounds as the signaling molecules antagonists [[100](#page-14-7)–[102](#page-14-8)]. Given that QS controls diferent phases of the bioflm development, including initial colonization/ adhesion, bacterial aggregation, bioflm maturation and cell dispersion, inhibiting QS will prevent the bioflm formation [\[103\]](#page-14-9). QSIs are, therefore, applied as a therapeutic agent in treating bioflm infection to inhibit the bioflm formation [\[104\]](#page-14-10). Antibiotics, synthetic compounds and natural products may infuence the QSIs function [\[99](#page-14-6)]. Some synthetic and natural compounds that act as QSIs and reduce the bioflm formation in bacteria are shown in Table [2.](#page-6-0) Drugs like aspirin, piroxicam and meloxicam, which are in the category of nonsteroidal anti-infammatory drugs, may be applied as potential inhibitors to control the QS signaling molecules of *P. aeruginosa*, as well as biofilm development [[105–](#page-14-11)[107](#page-14-12)]. Antibiotics such as azithromycin, erythromycin, ciprofoxacin, ceftazidime, gentamicin, tobramycin, piperacillin, spectinomycin and streptomycin display good levels of the QSI activity [\[108,](#page-14-13) [109\]](#page-14-14). Bacteria could be resistant to a single synthetic and natural compound, leading to decreasing the efective activity of them. Therefore, a combined therapy of antibiotics and QSIs is recommended. Consequently, this combined therapy can improve the efficiency of treatment without increasing the antibiotics toxicity and inhibiting the resistance to a single QSI [[110](#page-14-15), [111](#page-14-16)]. The combination of Aminoglycoside antibiotics with resveratrol signifcantly diminishes the bioflm production, as compared with each of the compounds alone  $[112]$  $[112]$ . The synergistic effectiveness of curcumin with ceftazidime, ciprofoxacin, gentamicin and azithromycin on the *P. aeruginosa* QS signaling molecule showed that the sub-MIC of each of the compounds, both alone and in combination, could signifcantly decline the bioflm development [[28,](#page-12-0) [113](#page-14-18)]. Zinc oxide (ZnO) nanoparticles, Chitosan and Chitosan-ZnO nanocomposite, in combination with gentamicin, signifcantly decreased the bioflm formation of both *S. aureus* and *P. aeruginosa*, when they were treated with MIC and 1/4 MIC of the compounds [[114](#page-14-19)]. Therefore, targeting QS by various anti-QS agents could be a potential application in the treatment of antibioflm infections.

#### **Aptamers**

Aptamers are peptides or single-stranded oligonucleotide molecules produced in an in vitro procedure called Systematic Evolution of Ligands by Exponential Enrichment (SELEX) [[141\]](#page-15-0). Due to their 3-dimensional constructions, they can be linked with great affinity and specificity to select target molecules including small molecules, proteins, drugs, metal ions, and even whole cells [[142](#page-15-1), [143](#page-15-2)]. Such properties of aptamers lead to a broad range of activities as antibioflm and antibacterial agents [\[144](#page-15-3)]. Bacterial cell-wall depolarization may be due to the antibacterial efects of aptamer [[145](#page-15-4)]. Although the available research is limited, several studies have suggested that aptamers could be used as an alternative strategy to inhibit the development of bioflms [[144](#page-15-3), [146](#page-15-5), [147\]](#page-15-6). Figure [2](#page-7-0) shows multiple mechanisms of aptamers for the suppression of the bacterial bioflm formation. The aim of these researches has been selection of aptamers against targets involved in the bioflm development. Thevendran et al. [[148](#page-15-7)] developed a quickly growing feld of research by scientists from various scientifc branches. The fexibility of aptamers as agents of both diagnostics and therapy has introduced them as good candidates for a broad range of uses. Bacterial fagella are responsible for the initial attachment and motility, which are necessary

Compounds	Biofilm inhibitor molecules	Bacteria affected	Ref
Natural chemicals	Halogenated Furanone	P. aeruginosa and Vibrio harveyi	[115, 116]
	Penicillium: penicillic acid and patulin	P. aeruginosa	[117]
	Ananas comosus	P. aeruginosa PAO1	$\lceil 118 \rceil$
	Grapefruit juice (furocoumarins)	E. coli	$[119]$
	Manilkara zapota	P. aeruginosa	$[118]$
	Ocimum sanctum	P. aeruginosa	$[118]$
	Medicinal plant extracts such as C. viminalis (leaves), C. erectus (leaves), B. buceras (leaves)	P. aeruginosa	$[120]$
	Combretum albiflorum (bark)	P. aeruginosa	[121]
	Eugenol from clove extract	P. aeruginosa	$[122]$
	Curcumin from Curcuma longa (turmeric)	P. aeruginosa, E. coli, Serratia marcescens and Proteus mirabilis	[123]
	Carvacrol	S. aureus and S. enterica subsp. typhimurium	[124]
	Diterpene phytol	P. aeruginosa	[125]
	6-Gingerol	P. aeruginosa	$[27]$
	Menthol from peppermint (Mentha piperita) oil	P. aeruginosa and Aeromonas hydrophila	[126]
	Zingerone	P. aeruginosa	[127]
	<b>Baicalin</b>	P. aeruginosa	$[128]$
	<b>Berberine</b>	P. aeruginosa	$\lceil 129 \rceil$
	Flavonoid fraction of Psidium guajava leaves	P. aeruginosa	[130]
	Ajoene from garlic	P. aeruginosa	$[131]$
	Diterpene phytol	P. aeruginosa	$[132]$
Synthetic chemicals	2-heptyl-6-nitro-4-oxo-1,4- dihydroquinoline-3-carboxamide	P. aeruginosa	[133]
	N-(indole-3-butanoyl)-L-HSL	P. aeruginosa PAO1	$[134]$
	Furanone C-30	P. aeruginosa	[135]
	<b>Brominated furanone</b>	S. anginosus, Staphylococcus intermedius and S. mutans	[136]
	Meta-bromo-thiolactone	P. aeruginosa	$\lceil 137 \rceil$
	MuPEP1 Inhibitors	S. mutans	$[138]$
	Pyridoxal lactohydrazone	P. aeruginosa	$[139]$
	Lacto hydrazone	P. aeruginosa	[140]

<span id="page-6-0"></span>**Table 2** Major recent developments in the inhibition of bioflm formation

for the bioflm development. Therefore, inhibiting fagella represents an efficient potential approach to prevent the bioflm development. Ning et al. [[142\]](#page-15-1) demonstrated that a specifc aptamer targeting the fagella of *Salmonella choleraesuis* could develop the antibioflms efect on the inhibition of the bacterial bioflms. Yu et al. [[149](#page-15-8)] also screened the aptamers targeted QS signal molecule that could interfere with the virulence gene expression of *P. aeruginosa*. Furthermore, Mao et al.  $[146]$  $[146]$  reported that aptamer-graphene oxide conjugates displayed excellent antibioflm properties against *Salmonella typhimurium* and could serve as a longterm approach to control the bacterial bioflm formation. In addition, Shatila et al. [[144\]](#page-15-3) demonstrated that the DNA aptamer (Apt17) targeted invasion protein A of *Salmonella*  *enteritidis* (SidA) could reduce the bioflm formation, either alone or in combination with ampicillin. Further, Wang et al. [\[150](#page-15-9)] revealed that the PA-ap1 aptamer in combination with SWNTs (Single-walled carbon nanotubes) could reduce the bioflm development of *P. aeruginosa* about 36%, in comparison to the SWNTs alone. In addition, the combination of ciprofoxacin with PA-ap1 aptamer–SWNTs had a greater antibiofilm efficacy than any compound alone. As bacteria accumulation represents a signifcant phase at the initial step of the bioflm development, aptamer-SWNTs could spontaneously catch and link to bacterial cells, quickly accumulating a greater efective concentration of aptamer-SWNTs around bacterial cells that could interfere with the bacteria accumulation. In addition, it was revealed that



<span id="page-7-0"></span>**Fig. 2** Multiple mechanisms of aptamers for the suppression of the bacterial bioflm formation

aptamer–SWNTs signifcantly decreased the bioflm formation, in comparison to the SWNTs alone, thus implying the possibility of using the targeted efective aptamers in the removal of the bioflm. Aptamers, as special targeting agents, can be used for the treatment of bacterial infections, which could increase the efficient concentration of antibiotics and decrease the off-target effects. It was reported that the combination of aptamer with antibiotics could lead to attacking more bacterial cells, in comparison to the use of antibiotics alone [\[142](#page-15-1)]. Wang et al. [\[150](#page-15-9)] also demonstrated that C4-HSL aptamers could efficiently decrease the biofilm development of *P. aeruginosa* with inhibiting QS, in comparison to that of the untreated groups. In enteropathogenic *E. coli* (EPEC), the bioflm formation is upregulated using some genes such as cell interaction (*lsrA*), motility (*motB*) and curli gene (*csgA*). Aptamer SELEX 10 Colony 5 could reduce the mRNA level of *csgA*, *lsrA* and *motB* gene, consequently decreasing the bioflm formation of the treated group

 $\mathcal{D}$  Springer

[\[151](#page-15-26)]. Sengupta et al. [\[152](#page-15-27)] also reported that aptamer-DNA templated Ag-NC (silver-nanocluster) could act as a sensor, thus creating a novel possibility to detect planktonic and bioflm forms. In addition, Ag-NC aptamer could be potentially efective in inhibiting the bioflm development in *P. aeruginosa*. Therefore, aptamer and aptamer coupled with excellent agents could provide high target specifcity to inhibit the bacterial growth and bioflm development; thus, they could be regarded as ideal strategies for the development of antibacterial and antibioflm agents.

#### **Nanoparticles**

Recently, nanoparticles have received much attention due to their antibacterial and antibioflm characteristics. Therefore, NPs can be presented as an alternative therapeutic strategy to inhibit the development of the bacterial bioflm [\[20\]](#page-11-19). Figure [3](#page-8-0) shows diferent physicochemical interactions between

<span id="page-8-0"></span>**Fig. 3** Diferent physicochemical interactions between bacterial bioflm and nanoparticles



bacterial bioflms and nanoparticles. One of the advantages of NPs is the high surface-area to volume ratio, which can provide a platform for the development of materials with a wide range of chemical, mechanical, magnetic and electrical characteristics [[153\]](#page-15-28). NPs can easily interact with the bacterial cell because of their shape, small size, surface charge, hydrophobicity and high surface-to-volume ratio [[154,](#page-15-29) [155](#page-15-30)].

NPs with a positive charge prefer to interact with the negatively charged components of the cell walls and plasma membranes of bacteria (polysaccharide, extracellular DNA (eDNA) and proteins). Furthermore, NPs with a negative charge prefer to interact with the positively charged components of the cell-wall, enhancing the bacterial membrane permeability and fowing out of the cytoplasm contents, which can cause the bacterial cell death. In addition, there are several water channels in the bacterial bioflm structure, which can enable bacteria to transport nutrients through these pores. Therefore, NPs can have an antibacterial efect on the bioflm through spreading from these channels [\[156](#page-16-0)].

The inhibition of the efflux pumps of bacteria by NPs is another potential mechanism for the antibioflm activity [\[157\]](#page-16-1). Gupta et al.  $[157]$  $[157]$  $[157]$  indicated that efflux pumps could play efective roles in the bacterial bioflm formation; as well, regulation of their expression could directly control the antibiotic resistance and development of bioflms. In addition, Padwal et al. [\[158\]](#page-16-2) showed the combination of antibiotics with NPs could inhibit the activity of efflux pumps, thus suggesting that these compounds could be applied as efflux pump inhibitors. They proposed that NPs could distract efflux kinetics through linking to the active site of efflux pumps and decreasing their activities, thus leading to

inhibition of the extrusion of antibiotics outside cells. Similar results have revealed the ZnO nanoparticles could have an inhibitory effect on such efflux pumps as MexAM-OPrM of *P. aeruginosa* and NorA of *S. aureus* [\[159](#page-16-3), [160\]](#page-16-4). Therefore, the downregulation or inhibition of efflux pumps with NPs could be regarded as a potential therapeutic approach to inhibit and/or decrease the bacterial bioflm development.

EPS produced by bacteria play critical roles in the primary attachment of the bacterial cells to the host cell surface and the development of a complex bioflm structure, which could protect bioflms against the classical antibiotics and host-immune system [\[161](#page-16-5), [162\]](#page-16-6). Production of EPS is one of the critical strategies to protect the bacterial bioflm against the antibiotic activity. Therefore, one of the mains restrictive strategies for the bioflm development is also EPS reduction [[25\]](#page-11-24). Inhibition of the EPS development can be one of the antibioflm mechanisms of NPs; it has been proved that NPs could reduce the bioflm development by the disruption of the EPS production [\[155](#page-15-30), [163\]](#page-16-7). In *S. aureus* bioflm, the chemical compounds of EPS contain polysaccharide, proteins and eDNA. One of the major compounds of the bioflm matrix is Polysaccharide Intercellular Adhesin (PIA), which is a compound with a positive charge. Cytoplasmic proteins and eDNA are positively and negatively charged, respectively [\[164](#page-16-8)]. NPs like chitosan NPs, due to having a positive charge, can bind to eDNA with a negative charge, disrupting the bioflm formation with their penetration into the bioflm structure [\[163\]](#page-16-7). In contrast, NPs with a negative charge like ZnO NPs can interact with positively charged PIA and modify the permeability of the bioflm structure; then they can easily penetrate into the bioflm

matrix, inhibiting the bioflm development by inducing ROS [\[164–](#page-16-8)[167\]](#page-16-9). The bioflm matrix of *P. aeruginosa* consists of exopolysaccharides and eDNA. In *P. aeruginosa*, there are three types of exopolysaccharides including alginate, Pel (a glucose-rich polysaccharide polymer encoded by *pel* operon including *pelA-F* genes), and Psl (composed of a repeating pentamer containing p-glucose, L-rhamnose, and p-mannose, which is encoded by the *psl* operon including *pslA-L*); they perform diferent roles in the bioflm development [[168,](#page-16-10) [169](#page-16-11)]. Alginate is a major compound of the bioflm matrix accounting for the cell surfaces binding and bioflm sustainability. Alginate and Pel are negatively and positively charged polysaccharides, respectively [\[170\]](#page-16-12). The positive and negative NPs could be charged by binding to alginate and Pel, respectively, which can prevent the attachment of the bacteria to cell surfaces; this is the early stage of bioflm development. These nanoparticles can disrupt the development of bioflm by preventing the bacterial attachment to the cellular surfaces [[163,](#page-16-7) [170](#page-16-12), [171\]](#page-16-13). Messiaen et al. [[172\]](#page-16-14) also demonstrated that the negatively charged components of tobramycin-loaded liposome were placed near the bacterial cell clusters of *Burkholderia cepacia* complex bioflms, while fber-like structures such as the extracellular DNA became immobilized by the interactions with the positively charged particles of nanospheres in the bioflm matrix.

In addition, owing to the antibacterial activity of NPs, they can reduce the bacterial attachment to surfaces and to themselves, as well as the replication and development of bioflm [\[154\]](#page-15-29). Table [3](#page-9-0) shows nanoparticles inhibiting the bacterial bioflm formation and development. According to these studies, NPs could be regarded as potential antibioflm agents as a function of their anti-adhesive activity, bactericidal activity and delivery capability. However, NPs may be toxic, but they could be modifed to reduce their toxicity, making them useful for biomedical applications. Therefore, further studies should be directed toward changing the oxidative state and charge density, by applying surface coatings and altering the ability to aggregate. Clinical and nonclinical studies are also needed to determine the safety and tolerance of NPs formulating the potential commercial applications.

#### **Peptide Nucleic Acids**

Peptide nucleic acids (PNAs) are structurally similar to DNA or RNA; they are the synthetic analogs of nucleic acid. Their structure is the repetition of *N*-(2-aminoethyl) glycine units in nucleotide bases instead of sugar phosphate back bones connected by unnatural pseudodipeptide bonds [\[182](#page-16-15)]. Properties of PNAs are a combination of both nucleic acids and peptides, which are chemically similar to protein and structurally similar to nucleic acid, which are relatively stable against enzymes degrading proteins and nucleic acids [[182,](#page-16-15) [183\]](#page-16-16). Antimicrobial and antibiofilm effects of PNAs are related to their small size and the targeted nucleotide

<span id="page-9-0"></span>**Table 3** Nanoparticles with the antibioflm properties

Nanoparticles	Mode of action	Bacteria affected	Refs.
FeOOH (iron-based NP)	Initial stage	P. aeruginosa	$\lceil 173 \rceil$
ZnO	Initial stage	P. aeruginosa, Streptococcus mitis and Streptococcus pneumoniae	[25, 74]
Silver	Initial stage	P. aeruginosa, S. aureus	[174, 175]
Gold	Initial stage	P. aeruginosa	[26]
Gentamicin-loaded gold	Maturation stage	S. aureus, E. coli, P. aeruginosa and Listeria monocytogenes	$[176]$
	Gentamicin-loaded nanotubes titanium Attachment disruption of the sessile cells to sur- faces (Initial stage)	S. aureus and S. epidermidis	$[177]$
Fucoidan-stabilized gold	Attachment disruption of the sessile cells to sur- faces (Initial stage)	P. aeruginosa	$[178]$
Crataeva nurvala		P. aeruginosa	$\lceil 179 \rceil$
Chitosan oligosaccharide-capped gold	Disrupted the attachment of sessile cells to surfaces (Initial stage)	P. aeruginosa	$\lceil 162 \rceil$
<b>Sulfonated Chitosan</b>	Secretion of exopolysaccharide and decreasing of the metabolic activity	S. <i>aureus</i> and <i>E. coli</i>	[180]
Titanium dioxide	Initial stage (inhibition EPS production)	S. mitis	$\lceil 181 \rceil$
Chlorhexidine-loaded mesoporous silica nanoparticles		Streptococcus sobrinus and S. mutans	[182]
Curcumin	Decrease of quorum sensing virulence factors	S. aureus and P. aeruginosa	[175]
Chitosan		S. aureus and P. aeruginosa	$\lceil 114 \rceil$
Chitosan-ZnO nanocomposite		S. aureus and P. aeruginosa	[114]

sequences of genes involved in the bioflm formation, in the absence or presence of a linker between the peptide conjugated to PNAs and its position in the structure of peptide-PNA  $[21]$ . The major property of PNA is its great affinity and specifcity binding. Therefore, it has a good potential to synthesize PNA-based specifc antibacterials for particular genes involved in the bioflm formation of the selected bacteria. PNAs demonstrate an enormous potential to inhibit the increase of the resistant bacteria [[184](#page-16-26)]. However, one of the most widespread negative properties of PNAs is the hydrophobicity of its structure, which could be due to its insolubility in the aqueous solution. The impermeability of bacterial membranes to PNA is another important limitation preventing the development of the antibacterial PNA due to the difculty in fnding the efective carriage and carriers of PNA to the bacterial cells [[183](#page-16-16)]. The lipid bilayer, lipopolysaccharide and peptidoglycan of bacteria serve as main barriers to the entrance of PNAs [[185](#page-16-27)].

Diferent mechanisms have been suggested to increase the transport of PNAs to bacterial cells: chemical changes of the PNAs structure can increase their hydrophilicity. In addition, covalent conjugation of various cell penetrating peptides (CPPs) with PNA [[186](#page-16-28)], such as photochemical internalization (PCI), chloroquine, cationic lipids and HIV-TAT [[187](#page-17-0)], and the use of the tetrahedral DNA nanostructure (TDN) can act a complementary base pair between PNAs and DNA [\[188\]](#page-17-1).

PNAs can prevent or overcome the bacterial biofilm development. PNAs treated at the *efaA* gene have a negative infuence on the bioflm development in *Enterococcus faecalis*; this gene serves an important role in the bacterial attachment to surfaces [[21](#page-11-20)]. In addition, Otsuka et al. [\[189\]](#page-17-2) showed that the *acpP* gene targeted by PNA-peptides caused antibacterial and antibioflm efects on both planktonic and bioflm states of *Haemophilus infuenzae*. Binding PNAs to their targeted genes could prevent and interfere with the gene expression. The *motA* gene could contribute to the downstream adhesion events of attachment in the development of bioflm in *P. aeruginosa,* playing a critical role in the initial state of the bioflm formation, which encodes the element of the fagellar motor complex [[190](#page-17-3)]. Xia et al. [\[191\]](#page-17-4) also revealed that PNA-peptides targeting the *motA* gene could decrease the *motA* expression and inhibit the bioflm development. PNAs have also been used in combination with classical antibiotics to increase their antibacterial and antibioflm activity. Castillo et al. [\[192\]](#page-17-5) showed the synergistic activity of mRNA targeted PNAs in *E. coli* O157:H7. The synergistic effect of trimethoprim and polymyxin B antibiotics in combination with anti-*acpP* PNA (encoding acyl carrier protein (AcpP) was determined using the checkerboard titration method. Their results showed that *acpP* mRNA targeted PNAs could improve the antibacterial activity of antibiotics by suppressing the *acpP* expression, which is functionally related with antibiotic resistance. In general, designing PNAs targeting specifc genes involved in the development and formation of bioflm, as well as targeting critical genes involved in the living and surviving bacteria, could enhance the susceptibility of bacteria and their bioflm.

# **Future Directions**

In the post-antibiotic era, development of novel antibioflm strategies and alternative therapeutic agents has been of interest to combat the threat of bacterial bioflm infections. In this era, we need more complex biological approaches to investigate the interactions between single- and multispecies bacterial bioflms and their environments in chronic infections. Some of the introduced strategies could only be used to treat single-species bioflms and may not be applicable to mixed-species bioflms. In addition, it is possible for bacteria to protect themselves during the use of these agents; this is a great challenge that should be addressed in regard to combating bacteria and their bioflm formation. Therefore, combination of these agents and fexibility in the use of them could be great promises for future bioflm infection treatments. The applicability of these strategies could be summarized in three steps including entry, delivery and release into the bioflm matrix. Recruitment of alternative and promising therapeutic approaches or strategies could increase the local concentration of antibiotics or antibioflm agents improve their local accumulation and decrease their systematic toxicity. Nanomedicine, as the most versatile and fexible strategy, can be combined with other novel therapeutic strategies as the state-of-art concepts. There are several tenuous antibioflm agents and a few of them could be used in clinical relevant models. Therefore, great efforts should be devoted to the evaluation of the biosafety and efficacy of strategies; effective routs for the entry, delivery and release of the agents to the bioflm matrix should be developed; further, the novel agents-bioflm interactions should be studies, and the complexity of prescription and cost of fabrication should be reduced.

## **Conclusions**

Antibioflm agents such as antiadhesion agents, antimicrobial peptides, aptamers, nanoparticles and peptide nucleic acids represent a promising therapeutic target to treat bioflm infections and develop a strong capability to interfere with diferent phases of the bioflm development, including adherence, polysaccharide intercellular adhesion (PIA), quorum sensing molecules and cell-to-cell connection, bacterial aggregation, planktonic bacteria killing and host-immune response modulation. Therefore, rational and experimental design approaches allow the synthesis of new antibioflm compounds with improved and varied biological functions. In addition, these novel strategies can be promising methods against bacterial bioflm due to their improved permeation and targeted delivery of antibacterials inside the bioflm. As well, these components in combination with antibiotics can lead to developing a powerful approach for the treatment of bacterial bioflm-related infections.

**Author Contributions** FH: Conceptualization; Data curation; Formal analysis; Investigation; Methodology; Roles/Writing—original draft; Writing—review & editing. MAR, SE, LY, RN & HSK: Data curation; Roles/Writing—original draft; Writing—review & editing. PG: Project administration; Resources; Software; Supervision; Validation; Visualization; Writing—review & editing.

# **Declarations**

**Conflict of interest** The authors declare that they have no confict of interest.

# **References**

- <span id="page-11-0"></span>1. Høiby, N., Bjarnsholt, T., Moser, C., Bassi, G., Coenye, T., Donelli, G., Hall-Stoodley, L., Holá, V., Imbert, C., & Kirketerp-Møller, K. (2015). ESCMID guideline for the diagnosis and treatment of bioflm infections 2014. *Clinical Microbiology & Infection, 21*, S1–S25.
- <span id="page-11-1"></span>2. Evans, J. J., & Bolz, D. D. (2019). Regulation of virulence and antibiotic resistance in Gram-positive microbes in response to cell wall-active antibiotics. *Current Opinion in Infectious Diseases, 32*, 217–222.
- <span id="page-11-2"></span>3. Khatoon, Z., McTiernan, C. D., Suuronen, E. J., Mah, T.-F., & Alarcon, E. I. (2018). Bacterial bioflm formation on implantable devices and approaches to its treatment and prevention. *Heliyon, 4*, e01067.
- <span id="page-11-3"></span>4. Batoni, G., Maisetta, G., & Esin, S. (2016). Antimicrobial peptides and their interaction with bioflms of medically relevant bacteria. *Biochimica et Biophysica Acta (BBA)-Biomembranes, 1858*, 1044–1060.
- <span id="page-11-4"></span>5. Lynch, A. S., & Robertson, G. T. (2008). Bacterial and fungal bioflm infections. *Annual Review of Medicine, 59*, 415–428.
- <span id="page-11-5"></span>6. Hall-Stoodley, L., Costerton, J. W., & Stoodley, P. (2004). Bacterial bioflms: From the natural environment to infectious diseases. *Nature Reviews Microbiology, 2*, 95–108.
- <span id="page-11-6"></span>7. Costerton, J. W., Cheng, K., Geesey, G. G., Ladd, T. I., Nickel, J. C., Dasgupta, M., & Marrie, T. J. (1987). Bacterial bioflms in nature and disease. *Annual Reviews in Microbiology, 41*, 435–464.
- <span id="page-11-7"></span>8. Lei, M. G., Gupta, R. K., & Lee, C. Y. (2017). Proteomics of *Staphylococcus aureus* bioflm matrix in a rat model of orthopedic implant-associated infection. *PLoS ONE, 12*, e0187981.
- <span id="page-11-8"></span>9. Watnick, P., & Kolter, R. (2000). Bioflm, city of microbes. *Journal of Bacteriology, 182*, 2675–2679.
- <span id="page-11-9"></span>10. Galdiero, E., Lombardi, L., Falanga, A., Libralato, G., Guida, M., & Carotenuto, R. (2019). Bioflms: Novel strategies based on antimicrobial peptides. *Pharmaceutics, 11*, 322.
- <span id="page-11-10"></span>11. Staudt, C., Horn, H., Hempel, D., & Neu, T. (2004). Volumetric measurements of bacterial cells and extracellular polymeric substance glycoconjugates in bioflms. *Biotechnology and Bioengineering, 88*, 585–592.
- <span id="page-11-11"></span>12. Dibdin, G. H., Assinder, S. J., Nichols, W. W., & Lambert, P. A. (1996). Mathematical model of β-lactam penetration into a bioflm of *Pseudomonas aeruginosa* while undergoing simultaneous inactivation by released β-lactamases. *Journal of Antimicrobial Chemotherapy, 38*, 757–769.
- <span id="page-11-12"></span>13. Bhagirath, A. Y., Li, Y., Somayajula, D., Dadashi, M., Badr, S., & Duan, K. (2016). Cystic fbrosis lung environment and *Pseudomonas aeruginosa* infection. *BMC Pulmonary Medicine, 16*, 1–22.
- <span id="page-11-13"></span>14. Ebbensgaard, A., Mordhorst, H., Overgaard, M. T., Nielsen, C. G., Aarestrup, F. M., & Hansen, E. B. (2015). Comparative evaluation of the antimicrobial activity of diferent antimicrobial peptides against a range of pathogenic bacteria. *PLoS ONE, 10*, e0144611.
- <span id="page-11-14"></span>15. Kåhrström, C. T. (2013). Entering a post-antibiotic era? *Nature Reviews Microbiology, 11*, 146–146.
- <span id="page-11-15"></span>16. Di Somma, A., Moretta, A., Canè, C., Cirillo, A., & Duilio, A. (2020). Antimicrobial and antibioflm peptides. *Biomolecules, 10*, 652.
- <span id="page-11-16"></span>17. Ferriol-González, C., & Domingo-Calap, P. (2020). Phages for bioflm removal. *Antibiotics, 9*, 268.
- <span id="page-11-17"></span>18. Haque, S., Ahmad, F., Dar, S. A., Jawed, A., Mandal, R. K., Wahid, M., Lohani, M., Khan, S., Singh, V., & Akhter, N. (2018). Developments in strategies for Quorum Sensing virulence factor inhibition to combat bacterial drug resistance. *Microbial Pathogenesis, 121*, 293–302.
- <span id="page-11-18"></span>19. Shatila, F., Yaşa, İ, & Yalçın, H. (2020). Inhibition of *Salmonella enteritidis* bioflms by Salmonella invasion protein-targeting aptamer. *Biotechnology Letters*. [https://doi.org/10.1007/](https://doi.org/10.1007/s10529-020-02920-2) [s10529-020-02920-2](https://doi.org/10.1007/s10529-020-02920-2)
- <span id="page-11-19"></span>20. Fulaz, S., Vitale, S., Quinn, L., & Casey, E. (2019). Nanoparticle–bioflm interactions: The role of the EPS matrix. *Trends in Microbiology, 27*, 915–926.
- <span id="page-11-20"></span>21. Narenji, H., Teymournejad, O., Rezaee, M. A., Taghizadeh, S., Mehramuz, B., Aghazadeh, M., Asgharzadeh, M., Madhi, M., Gholizadeh, P., Ganbarov, K., Yousef, M., Pakravan, A., Dal, T., Ahmadi, R., & Samadi Kafl, H. (2020). Antisense peptide nucleic acids againstftsZ andefaA genes inhibit growth and bioflm formation of *Enterococcus faecalis*. *Microbial Pathogenesis, 139*, 103907.
- <span id="page-11-21"></span>22. Arias, C. A., & Murray, B. E. (2009). Antibiotic-resistant bugs in the 21st century—A clinical super-challenge. *New England Journal of Medicine, 360*, 439–443.
- <span id="page-11-22"></span>23. Wood, L. F., Leech, A. J., & Ohman, D. E. (2006). Cell wallinhibitory antibiotics activate the alginate biosynthesis operon in *Pseudomonas aeruginosa*: roles of σ22 (AlgT) and the AlgW and Prc proteases. *Molecular Microbiology, 62*, 412–426.
- <span id="page-11-23"></span>24. Wiens, J. R., Vasil, A. I., Schurr, M. J., & Vasil, M. L. (2014). Iron-regulated expression of alginate production, mucoid phenotype, and bioflm formation by *Pseudomonas aeruginosa*. *MBio*. <https://doi.org/10.1128/mBio.01010-13>
- <span id="page-11-24"></span>25. Bhattacharyya, P., Agarwal, B., Goswami, M., Maiti, D., Baruah, S., & Tribedi, P. (2018). Zinc oxide nanoparticle inhibits the bioflm formation of *Streptococcus pneumoniae*. *Antonie van Leeuwenhoek, 111*, 89–99.
- <span id="page-11-26"></span>26. Ali, S. G., Ansari, M. A., Alzohairy, M. A., Alomary, M. N., AlYahya, S., Jalal, M., Khan, H. M., Asiri, S. M. M., Ahmad, W., & Mahdi, A. A. (2020). Biogenic gold nanoparticles as potent antibacterial and antibiofilm nano-antibiotics against *Pseudomonas aeruginosa*. *Antibiotics, 9*, 100.
- <span id="page-11-25"></span>27. Kim, H.-S., Lee, S.-H., Byun, Y., & Park, H.-D. (2015). 6-Gingerol reduces *Pseudomonas aeruginosa* bioflm formation and

virulence via quorum sensing inhibition. *Science and Reports, 5*, 8656.

- <span id="page-12-0"></span>28. Bahari, S., Zeighami, H., Mirshahabi, H., Roudashti, S., & Haghi, F. (2017). Inhibition of *Pseudomonas aeruginosa* quorum sensing by subinhibitory concentrations of curcumin with gentamicin and azithromycin. *Journal of Global Antimicrobial Resistance, 10*, 21–28.
- <span id="page-12-1"></span>29. De la Fuente-Núñez, C., Refuveille, F., Fernández, L., & Hancock, R. E. (2013). Bacterial bioflm development as a multicellular adaptation: Antibiotic resistance and new therapeutic strategies. *Current Opinion in Microbiology, 16*, 580–589.
- <span id="page-12-2"></span>30. Donlan, R. M. (2002). Bioflms: Microbial life on surfaces. *Emerging Infectious Diseases, 8*, 881.
- <span id="page-12-3"></span>31. Beloin, C., Valle, J., Latour-Lambert, P., Faure, P., Kzreminski, M., Balestrino, D., Haagensen, J. A., Molin, S., Prensier, G., & Arbeille, B. (2004). Global impact of mature bioflm lifestyle on *Escherichia coli* K-12 gene expression. *Molecular Microbiology, 51*, 659–674.
- <span id="page-12-4"></span>32. Klausen, M., Aaes-Jørgensen, A., Molin, S., & Tolker-Nielsen, T. (2003). Involvement of bacterial migration in the development of complex multicellular structures in *Pseudomonas aeruginosa* bioflms. *Molecular Microbiology, 50*, 61–68.
- 33. Klausen, M., Heydorn, A., Ragas, P., Lambertsen, L., Aaes-Jørgensen, A., Molin, S., & Tolker-Nielsen, T. (2003). Bioflm formation by *Pseudomonas aeruginosa* wild type, fagella and type IV pili mutants. *Molecular Microbiology, 48*, 1511–1524.
- 34. Toutain, C. M., Caizza, N. C., Zegans, M. E., & O'Toole, G. A. (2007). Roles for fagellar stators in bioflm formation by *Pseudomonas aeruginosa*. *Research in Microbiology, 158*, 471–477.
- <span id="page-12-5"></span>35. Watnick, P. I., & Kolter, R. (1999). Steps in the development of a *Vibrio cholerae* El Tor bioflm. *Molecular Microbiology, 34*, 586–595.
- <span id="page-12-6"></span>36. Schmidt, J., Müsken, M., Becker, T., Magnowska, Z., Bertinetti, D., Möller, S., Zimmermann, B., Herberg, F. W., Jänsch, L., & Häussler, S. (2011). The *Pseudomonas aeruginosa* chemotaxis methyltransferase CheR1 impacts on bacterial surface sampling. *PLoS ONE, 6*, e18184.
- <span id="page-12-7"></span>37. Hadjifrangiskou, M., Gu, A. P., Pinkner, J. S., Kostakioti, M., Zhang, E. W., Greene, S. E., & Hultgren, S. J. (2012). Transposon mutagenesis identifes uropathogenic *Escherichia coli* bioflm factors. *Journal of Bacteriology, 194*, 6195–6205.
- <span id="page-12-8"></span>38. Beloin, C., Roux, A., & Ghigo, J.-M. (2008). *Escherichia coli* bioflms. In *Bacterial bioflms.* (pp. 249–289). Springer.
- <span id="page-12-9"></span>39. Anderson, G. G., Palermo, J. J., Schilling, J. D., Roth, R., Heuser, J., & Hultgren, S. J. (2003). Intracellular bacterial bioflm-like pods in urinary tract infections. *Science, 301*, 105–107.
- <span id="page-12-10"></span>40. Waksman, G., & Hultgren, S. J. (2009). Structural biology of the chaperone–usher pathway of pilus biogenesis. *Nature Reviews Microbiology, 7*, 765–774.
- <span id="page-12-11"></span>41. Spurbeck, R. R., Stapleton, A. E., Johnson, J. R., Walk, S. T., Hooton, T. M., & Mobley, H. L. (2011). Fimbrial profles predict virulence of uropathogenic *Escherichia coli* strains: contribution of ygi and yad fmbriae. *Infection and Immunity, 79*, 4753–4763.
- <span id="page-12-12"></span>42. Thumbikat, P., Berry, R. E., Zhou, G., Billips, B. K., Yaggie, R. E., Zaichuk, T., Sun, T.-T., Schaefer, A. J., & Klumpp, D. J. (2009). Bacteria-induced uroplakin signaling mediates bladder response to infection. *PLoS Pathogens, 5*, e1000415.
- <span id="page-12-13"></span>43. Cegelski, L., Pinkner, J. S., Hammer, N. D., Cusumano, C. K., Hung, C. S., Chorell, E., Åberg, V., Walker, J. N., Seed, P. C., & Almqvist, F. (2009). Small-molecule inhibitors target *Escherichia coli* amyloid biogenesis and bioflm formation. *Nature Chemical Biology, 5*, 913–919.
- <span id="page-12-14"></span>44. Klebensberger, J., Birkenmaier, A., Gefers, R., Kjelleberg, S., & Philipp, B. (2009). SiaA and SiaD are essential for inducing autoaggregation as a specifc response to detergent stress in

*Pseudomonas aeruginosa*. *Environmental Microbiology, 11*, 3073–3086.

- <span id="page-12-15"></span>45. Najafi, K., Ganbarov, K., Gholizadeh, P., Tanomand, A., Rezaee, M. A., Mahmood, S. S., Asgharzadeh, M., & Kafl, H. S. (2020). Oral cavity infection by *Enterococcus faecalis*: Virulence factors and pathogenesis. *Reviews in Medical Microbiology, 31*, 51–60.
- <span id="page-12-16"></span>46. Sillanpää, J., Nallapareddy, S. R., Prakash, V. P., Qin, X., Hook, M., Weinstock, G. M., & Murray, B. E. (2008). Identifcation and phenotypic characterization of a second collagen adhesin, Scm, and genome-based identifcation and analysis of 13 other predicted MSCRAMMs, including four distinct pilus loci, in *Enterococcus faecium*. *Microbiology (Reading, England), 154*, 3199.
- <span id="page-12-17"></span>47. Han, Z., Pinkner, J. S., Ford, B., Obermann, R., Nolan, W., Wildman, S. A., Hobbs, D., Ellenberger, T., Cusumano, C. K., & Hultgren, S. J. (2010). Structure-based drug design and optimization of mannoside bacterial FimH antagonists. *Journal of Medicinal Chemistry, 53*, 4779–4792.
- <span id="page-12-18"></span>48. Bouckaert, J., Berglund, J., Schembri, M., De Genst, E., Cools, L., Wuhrer, M., Hung, C. S., Pinkner, J., Slättegård, R., & Zavialov, A. (2005). Receptor binding studies disclose a novel class of high-afnity inhibitors of the *Escherichia coli* FimH adhesin. *Molecular Microbiology, 55*, 441–455.
- <span id="page-12-19"></span>49. Cusumano, C. K., Pinkner, J. S., Han, Z., Greene, S. E., Ford, B. A., Crowley, J. R., Henderson, J. P., Janetka, J. W., & Hultgren, S. J. (2011). Treatment and prevention of urinary tract infection with orally active FimH inhibitors. *Science Translational Medicine, 3*, 109ra115.
- <span id="page-12-20"></span>50. Chorell, E., Pinkner, J. S., Phan, G., Edvinsson, S., Buelens, F., Remaut, H., Waksman, G., Hultgren, S. J., & Almqvist, F. (2010). Design and synthesis of C-2 substituted thiazolo and dihydrothiazolo ring-fused 2-pyridones: Pilicides with increased antivirulence activity. *Journal of Medicinal Chemistry, 53*, 5690–5695.
- <span id="page-12-21"></span>51. Chorell, E., Bengtsson, C., Banchelin, T.S.-L., Das, P., Uvell, H., Sinha, A. K., Pinkner, J. S., Hultgren, S. J., & Almqvist, F. (2011). Synthesis and application of a bromomethyl substituted scaffold to be used for efficient optimization of antivirulence activity. *European Journal of Medicinal Chemistry, 46*, 1103–1116.
- <span id="page-12-22"></span>52. Yazici, A., Ortucu, S., Taskin, M., & Marinelli, L. (2018). Natural-based antibioflm and antimicrobial peptides from micro-organisms. *Current Topics in Medicinal Chemistry, 18*, 2102–2107.
- <span id="page-12-23"></span>53. Ageitos, J., Sánchez-Pérez, A., Calo-Mata, P., & Villa, T. (2017). Antimicrobial peptides (AMPs): Ancient compounds that represent novel weapons in the fght against bacteria. *Biochemical Pharmacology, 133*, 117–138.
- <span id="page-12-24"></span>54. Zhang, L., & Falla, T. J. (2006). Antimicrobial peptides: Therapeutic potential. *Expert Opinion on Pharmacotherapy, 7*, 653–663.
- <span id="page-12-25"></span>55. Yeaman, M. R., & Yount, N. Y. (2003). Mechanisms of antimicrobial peptide action and resistance. *Pharmacological Reviews, 55*, 27–55.
- <span id="page-12-26"></span>56. Jenssen, H., Hamill, P., & Hancock, R. E. (2006). Peptide antimicrobial agents. *Clinical Microbiology Reviews, 19*, 491–511.
- <span id="page-12-27"></span>57. Brogden, K. A. (2005). Antimicrobial peptides: Pore formers or metabolic inhibitors in bacteria? *Nature Reviews Microbiology, 3*, 238–250.
- 58. Cassone, M., Frith, N., Vogiatzi, P., Wade, J. D., & Otvos, L. (2009). Induced resistance to the designer proline-rich antimicrobial peptide A3-APO does not involve changes in the intracellular target DnaK. *International Journal of Peptide Research and Therapeutics, 15*, 121–128.
- 59. Shah, P., Hsiao, F. S. H., Ho, Y. H., & Chen, C. S. (2016). The proteome targets of intracellular targeting antimicrobial peptides. *Proteomics, 16*, 1225–1237.
- <span id="page-13-10"></span>60. Graf, M., & Wilson, D. N. (2019). Intracellular antimicrobial peptides targeting the protein synthesis machinery. In *Antimicrobial peptides.* (pp. 73–89). Springer.
- <span id="page-13-0"></span>61. Okuda, K.-I., Zendo, T., Sugimoto, S., Iwase, T., Tajima, A., Yamada, S., Sonomoto, K., & Mizunoe, Y. (2013). Efects of bacteriocins on methicillin-resistant *Staphylococcus aureus* bioflm. *Antimicrobial Agents and Chemotherapy, 57*, 5572–5579.
- <span id="page-13-1"></span>62. Overhage, J., Campisano, A., Bains, M., Torfs, E. C., Rehm, B. H., & Hancock, R. E. (2008). Human host defense peptide LL-37 prevents bacterial bioflm formation. *Infection and Immunity, 76*, 4176–4182.
- <span id="page-13-2"></span>63. Brancatisano, F. L., Maisetta, G., Di Luca, M., Esin, S., Bottai, D., Bizzarri, R., Campa, M., & Batoni, G. (2014). Inhibitory efect of the human liver-derived antimicrobial peptide hepcidin 20 on bioflms of polysaccharide intercellular adhesin (PIA) positive and PIA-negative strains of *Staphylococcus epidermidis*. *Biofouling, 30*, 435–446.
- <span id="page-13-3"></span>64. Blower, R. J., Barksdale, S. M., & van Hoek, M. L. (2015). Snake cathelicidin NA-CATH and smaller helical antimicrobial peptides are efective against *Burkholderia thailandensis*. *PLoS Neglected Tropical Diseases, 9*, e0003862.
- <span id="page-13-4"></span>65. Anunthawan, T., De La Fuente-Núñez, C., Hancock, R. E., & Klaynongsruang, S. (2015). Cationic amphipathic peptides KT2 and RT2 are taken up into bacterial cells and kill planktonic and bioflm bacteria. *Biochimica et Biophysica Acta (BBA)-Biomembranes, 1848*, 1352–1358.
- <span id="page-13-5"></span>66. De Brucker, K., Delattin, N., Robijns, S., Steenackers, H., Verstraeten, N., Landuyt, B., Luyten, W., Schoofs, L., Dovgan, B., & Fröhlich, M. (2014). Derivatives of the mouse cathelicidinrelated antimicrobial peptide (CRAMP) inhibit fungal and bacterial bioflm formation. *Antimicrobial Agents and Chemotherapy, 58*, 5395–5404.
- <span id="page-13-6"></span>67. Mataraci, E., & Dosler, S. (2012). In vitro activities of antibiotics and antimicrobial cationic peptides alone and in combination against methicillin-resistant *Staphylococcus aureus* bioflms. *Antimicrobial Agents and Chemotherapy, 56*, 6366–6371.
- <span id="page-13-7"></span>68. Cao, Y., Yin, H., Wang, W., Pei, P., Wang, Y., Wang, X., Jiang, J., Luo, S.-Z., & Chen, L. (2020). Killing *Streptococcus mutans* in mature bioflm with a combination of antimicrobial and antibioflm peptides. *Amino Acids, 52*, 1–14.
- <span id="page-13-8"></span>69. Gopal, R., Lee, J. H., Kim, Y. G., Kim, M.-S., Seo, C. H., & Park, Y. (2013). Anti-microbial, anti-bioflm activities and cell selectivity of the NRC-16 peptide derived from witch flounder, *Glyptocephalus cynoglossus*. *Marine Drugs, 11*, 1836–1852.
- <span id="page-13-9"></span>70. Chen, L., Jia, L., Zhang, Q., Zhou, X., Liu, Z., Li, B., Zhu, Z., Wang, F., Yu, C., & Zhang, Q. (2017). A novel antimicrobial peptide against dental-caries-associated bacteria. *Anaerobe, 47*, 165–172.
- <span id="page-13-11"></span>71. Cooper, V. S., Carlson, W. A., & LiPuma, J. J. (2009). Susceptibility of *Caenorhabditis elegans* to Burkholderia infection depends on prior diet and secreted bacterial attractants. *PLoS ONE, 4*, e7961.
- <span id="page-13-12"></span>72. Peng, X., Zhang, Y., Bai, G., Zhou, X., & Wu, H. (2016). Cyclic di-AMP mediates bioflm formation. *Molecular Microbiology, 99*, 945–959.
- <span id="page-13-13"></span>73. Pletzer, D., Coleman, S. R., & Hancock, R. E. (2016). Anti-bioflm peptides as a new weapon in antimicrobial warfare. *Current Opinion in Microbiology, 33*, 35–40.
- <span id="page-13-14"></span>74. Ribeiro, S. M., De La Fuente-Núñez, C., Baquir, B., Faria-Junior, C., Franco, O. L., & Hancock, R. E. (2015). Antibioflm peptides increase the susceptibility of carbapenemase-producing *Klebsiella pneumoniae* clinical isolates to β-lactam antibiotics. *Antimicrobial Agents and Chemotherapy, 59*, 3906–3912.
- <span id="page-13-15"></span>75. Tong, Z., Zhang, Y., Ling, J., Ma, J., Huang, L., & Zhang, L. (2014). An in vitro study on the efects of nisin on the antibacterial activities of 18 antibiotics against *Enterococcus faecalis*. *PLoS ONE, 9*, e89209.
- <span id="page-13-16"></span>76. Tong, Z., Zhou, L., Jiang, W., Kuang, R., Li, J., Tao, R., & Ni, L. (2011). An in vitro synergetic evaluation of the use of nisin and sodium fuoride or chlorhexidine against *Streptococcus mutans*. *Peptides, 32*, 2021–2026.
- <span id="page-13-17"></span>77. Domingo-Calap, P., & Delgado-Martínez, J. (2018). Bacteriophages: Protagonists of a post-antibiotic era. *Antibiotics, 7*, 66.
- <span id="page-13-18"></span>78. Principi, N., Silvestri, E., & Esposito, S. (2019). Advantages and limitations of bacteriophages for the treatment of bacterial infections. *Frontiers in Pharmacology, 10*, 513.
- <span id="page-13-19"></span>79. Bedi, M. S., Verma, V., & Chhibber, S. (2009). Amoxicillin and specifc bacteriophage can be used together for eradication of bioflm of *Klebsiella pneumoniae* B5055. *World Journal of Microbiology & Biotechnology, 25*, 1145.
- <span id="page-13-20"></span>80. Lu, T. K., & Collins, J. J. (2007). Dispersing bioflms with engineered enzymatic bacteriophage. *Proceedings of the National Academy of Sciences, 104*, 11197–11202.
- <span id="page-13-21"></span>81. Łusiak-Szelachowska, M., Weber-Dąbrowska, B., & Górski, A. (2020). Bacteriophages and lysins in bioflm control. *Virologica Sinica*.<https://doi.org/10.1007/s12250-019-00192-3>
- <span id="page-13-22"></span>82. Wood, S., Kirkham, J., Marsh, P., Shore, R., Nattress, B., & Robinson, C. (2000). Architecture of intact natural human plaque bioflms studied by confocal laser scanning microscopy. *Journal of Dental Research, 79*, 21–27.
- <span id="page-13-23"></span>83. Pires, D. P., Oliveira, H., Melo, L. D., Sillankorva, S., & Azeredo, J. (2016). Bacteriophage-encoded depolymerases: Their diversity and biotechnological applications. *Applied Microbiology and Biotechnology, 100*, 2141–2151.
- <span id="page-13-24"></span>84. Cornelissen, A., Ceyssens, P. J., T'syen, J., Van Praet, H., Noben, J. P., Shaburova, O. V., Krylov, V. N., Volckaert, G., & Lavigne, R. (2011). The T7-related *Pseudomonas putida* phage φ15 displays virion-associated bioflm degradation properties. *PLoS ONE, 6*, e18597.
- <span id="page-13-25"></span>85. Yoon, S., Choi, Y., Lee, S. Y., Son, J., Jun, S., & Kang, S. (2013) Bacteriophage or lytic protein derived from the bacteriophage which efective for the treatment of *Staphylococcus aureus* bioflm, Google Patents.
- <span id="page-13-26"></span>86. Borysowski, J., Łobocka, M., Międzybrodzki, R., Weber-Dąbrowska, B., & Górski, A. (2011). Potential of bacteriophages and their lysins in the treatment of MRSA. *BioDrugs, 25*, 347–355.
- <span id="page-13-27"></span>87. Fischetti, V. A. (2018). Development of phage lysins as novel therapeutics: A historical perspective. *Viruses, 10*, 310.
- <span id="page-13-28"></span>88. Sharma, U., Vipra, A., & Channabasappa, S. (2018). Phagederived lysins as potential agents for eradicating bioflms and persisters. *Drug Discovery Today, 23*, 848–856.
- <span id="page-13-29"></span>89. Gray, J. A., Chandry, P. S., Kaur, M., Kocharunchitt, C., Bowman, J. P., & Fox, E. M. (2018). Novel biocontrol methods for *Listeria monocytogenes* bioflms in food production facilities. *Frontiers in Microbiology, 9*, 605.
- <span id="page-13-30"></span>90. Briers, Y., Walmagh, M., Grymonprez, B., Biebl, M., Pirnay, J.-P., Defraine, V., Michiels, J., Cenens, W., Aertsen, A., & Miller, S.  $(2014)$ . Art-175 is a highly efficient antibacterial against multidrug-resistant strains and persisters of *Pseudomonas aeruginosa*. *Antimicrobial Agents and Chemotherapy, 58*, 3774–3784.
- <span id="page-13-31"></span>91. Donlan, R. M. (2009). Preventing bioflms of clinically relevant organisms using bacteriophage. *Trends in Microbiology, 17*, 66–72.
- <span id="page-13-32"></span>92. Coulter, L. B., McLean, R. J., Rohde, R. E., & Aron, G. M. (2014). Efect of bacteriophage infection in combination with tobramycin on the emergence of resistance in *Escherichia coli* and *Pseudomonas aeruginosa* bioflms. *Viruses, 6*, 3778–3786.
- <span id="page-14-0"></span>93. Issa, R., Chanishvili, N., Caplin, J., Kakabadze, E., Bakuradze, N., Makalatia, K., & Cooper, I. (2019). Antibioflm potential of purifed environmental bacteriophage preparations against early stage *Pseudomonas aeruginosa* bioflms. *Journal of Applied Microbiology, 126*, 1657–1667.
- <span id="page-14-1"></span>94. Maciejewska, B., Olszak, T., & Drulis-Kawa, Z. (2018). Applications of bacteriophages versus phage enzymes to combat and cure bacterial infections: An ambitious and also a realistic application? *Applied Microbiology and Biotechnology, 102*, 2563–2581.
- <span id="page-14-2"></span>95. Abedon, S. T. (2019). Phage-antibiotic combination treatments: Antagonistic impacts of antibiotics on the pharmacodynamics of phage therapy? *Antibiotics, 8*, 182.
- <span id="page-14-3"></span>96. Tagliaferri, T. L., Jansen, M., & Horz, H.-P. (2019). Fighting pathogenic bacteria on two fronts: Phages and antibiotics as combined strategy. *Frontiers in Cellular and Infection Microbiology, 9*, 22.
- <span id="page-14-4"></span>97. Fuqua, C., Parsek, M. R., & Greenberg, E. P. (2001). Regulation of gene expression by cell-to-cell communication: Acylhomoserine lactone quorum sensing. *Annual Review of Genetics, 35*, 439–468.
- <span id="page-14-5"></span>98. Dijkshoorn, L., Nemec, A., & Seifert, H. (2007). An increasing threat in hospitals: Multidrug-resistant *Acinetobacter baumannii*. *Nature Reviews Microbiology, 5*, 939–951.
- <span id="page-14-6"></span>99. Hemmati, F., Salehi, R., Ghotaslou, R., Kafl, H. S., Hasani, A., Gholizadeh, P., Nouri, R., & Rezaee, M. A. (2020). Quorum Quenching: A potential target for antipseudomonal therapy. *Infection and Drug Resistance, 13*, 2989–3005.
- <span id="page-14-7"></span>100. Kalia, V. C. (2013). Quorum sensing inhibitors: An overview. *Biotechnology Advances, 31*, 224–245.
- 101. Dembitsky, V. M., Al Quntar, A. A. A., & Srebnik, M. (2011). Natural and synthetic small boron-containing molecules as potential inhibitors of bacterial and fungal quorum sensing. *Chemical Reviews, 111*, 209–237.
- <span id="page-14-8"></span>102. Kaufmann, G. F., Sartorio, R., Lee, S.-H., Mee, J. M., Altobell, L. J., Kujawa, D. P., Jefries, E., Clapham, B., Meijler, M. M., & Janda, K. D. (2006). Antibody interference with N-acyl homoserine lactone-mediated bacterial quorum sensing. *Journal of the American Chemical Society, 128*, 2802–2803.
- <span id="page-14-9"></span>103. Bassler, B. L. (2002). Small talk: Cell-to-cell communication in bacteria. *Cell, 109*, 421–424.
- <span id="page-14-10"></span>104. Taylor, P. K., Yeung, A. T., & Hancock, R. E. (2014). Antibiotic resistance in *Pseudomonas aeruginosa* bioflms: Towards the development of novel anti-bioflm therapies. *Journal of Biotechnology, 191*, 121–130.
- <span id="page-14-11"></span>105. de Almeida, F. A., Vargas, E. L. G., Carneiro, D. G., Pinto, U. M., & Vanetti, M. C. D. (2018). Virtual screening of plant compounds and nonsteroidal anti-infammatory drugs for inhibition of quorum sensing and bioflm formation in Salmonella. *Microbial Pathogenesis, 121*, 369–388.
- 106. Yang, E., Lu, Y., Xu, Y., Liang, Q., Wang, C., Wang, H., & Shen, H. (2014). Recombinant BCG coexpressing Ag85B, ESAT-6 and Rv3620c elicits specifc Th1 immune responses in C57BL/6 mice. *Microbial Pathogenesis, 69*, 53–59.
- <span id="page-14-12"></span>107. Soheili, V., Bazzaz, B. S. F., Abdollahpour, N., & Hadizadeh, F. (2015). Investigation of *Pseudomonas aeruginosa* quorumsensing signaling system for identifying multiple inhibitors using molecular docking and structural analysis methodology. *Microbial Pathogenesis, 89*, 73–78.
- <span id="page-14-13"></span>108. Skindersoe, M. E., Alhede, M., Phipps, R., Yang, L., Jensen, P. O., Rasmussen, T. B., Bjarnsholt, T., Tolker-Nielsen, T., Høiby, N., & Givskov, M. (2008). Efects of antibiotics on quorum sensing in *Pseudomonas aeruginosa*. *Antimicrobial Agents and Chemotherapy, 52*, 3648–3663.
- <span id="page-14-14"></span>109. Sofer, D., Gilboa-Garber, N., Belz, A., & Garber, N. C. (1999). 'Subinhibitory'erythromycin represses production of

*Pseudomonas aeruginosa* lectins, autoinducer and virulence factors. *Chemotherapy, 45*, 335–341.

- <span id="page-14-15"></span>110. Dong, Y.-H., & Zhang, L.-H. (2005). Quorum sensing and quorum-quenching enzymes. *The Journal of Microbiology, 43*, 101–109.
- <span id="page-14-16"></span>111. Chanda, S., & Rakholiya, K. (2011). Combination therapy: Synergism between natural plant extracts and antibiotics against infectious diseases. *Microbiology Book Series, 1*, 520–529.
- <span id="page-14-17"></span>112. Zhou, J.-W., Chen, T.-T., Tan, X.-J., Sheng, J.-Y., & Jia, A.-Q. (2018). Can the quorum sensing inhibitor resveratrol function as an aminoglycoside antibiotic accelerant against *Pseudomonas aeruginosa*? *International Journal of Antimicrobial Agents, 52*, 35–41.
- <span id="page-14-18"></span>113. Roudashti, S., Zeighami, H., Mirshahabi, H., Bahari, S., Soltani, A., & Haghi, F. (2017). Synergistic activity of sub-inhibitory concentrations of curcumin with ceftazidime and ciprofoxacin against *Pseudomonas aeruginosa* quorum sensing related genes and virulence traits. *World Journal of Microbiology & Biotechnology, 33*, 50.
- <span id="page-14-19"></span>114. Hemmati, F., Salehi, R., Ghotaslou, R., Kafl, H. S., Hasani, A., Gholizadeh, P., & Rezaee, M. A. (2020). The assessment of antibioflm activity of chitosan-zinc oxide-gentamicin nanocomposite on *Pseudomonas aeruginosa* and *Staphylococcus aureus*. *International Journal of Biological Macromolecules, 163*, 2248–2258.
- <span id="page-14-20"></span>115. de Nys, R., Wright, A. D., König, G. M., & Sticher, O. (1993). New halogenated furanones from the marine alga *Delisea pulchra* (cf. fmbriata). *Tetrahedron, 49*, 11213–11220.
- <span id="page-14-21"></span>116. De Nys, R., Givskov, M., Kumar, N., Kjelleberg, S., & Steinberg, P. (2006). Furanones. In *Antifouling compounds.* (pp. 55–86). Springer.
- <span id="page-14-22"></span>117. Rasmussen, T. B., Skindersoe, M. E., Bjarnsholt, T., Phipps, R. K., Christensen, K. B., Jensen, P. O., Andersen, J. B., Koch, B., Larsen, T. O., & Hentzer, M. (2005). Identity and efects of quorum-sensing inhibitors produced by Penicillium species. *Microbiology, 151*, 1325–1340.
- <span id="page-14-23"></span>118. Musthafa, K. S., Ravi, A. V., Annapoorani, A., Packiavathy, I. S. V., & Pandian, S. K. (2010). Evaluation of anti-quorumsensing activity of edible plants and fruits through inhibition of the N-acyl-homoserine lactone system in *Chromobacterium violaceum* and *Pseudomonas aeruginosa*. *Chemotherapy, 56*, 333–339.
- <span id="page-14-24"></span>119. Girennavar, B., Cepeda, M. L., Soni, K. A., Vikram, A., Jesudhasan, P., Jayaprakasha, G., Pillai, S. D., & Patil, B. S. (2008). Grapefruit juice and its furocoumarins inhibits autoinducer signaling and bioflm formation in bacteria. *International Journal of Food Microbiology, 125*, 204–208.
- <span id="page-14-25"></span>120. Adonizio, A., Kong, K.-F., & Mathee, K. (2008). Inhibition of quorum sensing-controlled virulence factor production in *Pseudomonas aeruginosa* by South Florida plant extracts. *Antimicrobial Agents and Chemotherapy, 52*, 198–203.
- <span id="page-14-26"></span>121. Vandeputte, O. M., Kiendrebeogo, M., Rajaonson, S., Diallo, B., Mol, A., El Jaziri, M., & Baucher, M. (2010). Identifcation of catechin as one of the favonoids from *Combretum albiforum* bark extract that reduces the production of quorum-sensingcontrolled virulence factors in *Pseudomonas aeruginosa* PAO1. *Applied and Environment Microbiology, 76*, 243–253.
- <span id="page-14-27"></span>122. Zhou, L., Zheng, H., Tang, Y., Yu, W., & Gong, Q. (2013). Eugenol inhibits quorum sensing at sub-inhibitory concentrations. *Biotechnology Letters, 35*, 631–637.
- <span id="page-14-28"></span>123. Packiavathy, I. A. S. V., Priya, S., Pandian, S. K., & Ravi, A. V. (2014). Inhibition of bioflm development of uropathogens by curcumin—An anti-quorum sensing agent from *Curcuma longa*. *Food Chemistry, 148*, 453–460.
- <span id="page-14-29"></span>124. Burt, S. A., Ojo-Fakunle, V. T., Woertman, J., & Veldhuizen, E. J. (2014). The natural antimicrobial carvacrol inhibits quorum

sensing in *Chromobacterium violaceum* and reduces bacterial bioflm formation at sub-lethal concentrations. *PLoS ONE, 9*, e93414.

- <span id="page-15-10"></span>125. Pejin, B., Ciric, A., Glamoclija, J., Nikolic, M., & Sokovic, M. (2015). In vitro anti-quorum sensing activity of phytol. *Natural Product Research, 29*, 374–377.
- <span id="page-15-11"></span>126. Husain, F. M., Ahmad, I., Khan, M. S., Ahmad, E., Tahseen, Q., Khan, M. S., & Alshabib, N. A. (2015). Sub-MICs of *Mentha piperita* essential oil and menthol inhibits AHL mediated quorum sensing and bioflm of Gram-negative bacteria. *Frontiers in Microbiology, 6*, 420.
- <span id="page-15-12"></span>127. Kumar, L., Chhibber, S., Kumar, R., Kumar, M., & Harjai, K. (2015). Zingerone silences quorum sensing and attenuates virulence of *Pseudomonas aeruginosa*. *Fitoterapia, 102*, 84–95.
- <span id="page-15-13"></span>128. Luo, J., Dong, B., Wang, K., Cai, S., Liu, T., Cheng, X., Lei, D., Chen, Y., Li, Y., & Kong, J. (2017). Baicalin inhibits bioflm formation, attenuates the quorum sensing-controlled virulence and enhances *Pseudomonas aeruginosa* clearance in a mouse peritoneal implant infection model. *PLoS ONE, 12*, e0176883.
- <span id="page-15-14"></span>129. Li, Y., Huang, J., Li, L., & Liu, L. (2017). Synergistic activity of berberine with azithromycin against *Pseudomonas aeruginosa* isolated from patients with cystic fbrosis of lung in vitro and in vivo. *Cellular Physiology and Biochemistry, 42*, 1657–1669.
- <span id="page-15-15"></span>130. Vasavi, H. S., Arun, A. B., & Rekha, P. D. (2014). Anti-quorum sensing activity of *Psidium guajava* L. favonoids against *Chromobacterium violaceum* and *Pseudomonas aeruginosa* PAO1. *Microbiology and Immunology, 58*, 286–293.
- <span id="page-15-16"></span>131. Jakobsen, T. H., Warming, A. N., Vejborg, R. M., Moscoso, J. A., Stegger, M., Lorenzen, F., Rybtke, M., Andersen, J. B., Petersen, R., & Andersen, P. S. (2017). A broad range quorum sensing inhibitor working through sRNA inhibition. *Science and Reports, 7*, 1–12.
- <span id="page-15-17"></span>132. Ilic-Tomić, T., Soković, M., Vojnović, S., Ćirić, A. D., Veljić, M., Nikodinović-Runić, J., & Novaković, M. M. (2017). Diarylheptanoids from *Alnus viridis* ssp viridis and *Alnus glutinosa*: Modulation of quorum sensing activity in *Pseudomonas aeruginosa*. *Planta Medica, 83*, 117–125.
- <span id="page-15-18"></span>133. Nafee, N., Husari, A., Maurer, C. K., Lu, C., de Rossi, C., Steinbach, A., Hartmann, R. W., Lehr, C.-M., & Schneider, M. (2014). Antibiotic-free nanotherapeutics: Ultra-small, mucus-penetrating solid lipid nanoparticles enhance the pulmonary delivery and anti-virulence efficacy of novel quorum sensing inhibitors. *Journal of Controlled Release, 192*, 131–140.
- <span id="page-15-19"></span>134. Geske, G. D., Wezeman, R. J., Siegel, A. P., & Blackwell, H. E. (2005). Small molecule inhibitors of bacterial quorum sensing and bioflm formation. *Journal of the American Chemical Society, 127*, 12762–12763.
- <span id="page-15-20"></span>135. Hentzer, M., Riedel, K., Rasmussen, T. B., Heydorn, A., Andersen, J. B., Parsek, M. R., Rice, S. A., Eberl, L., Molin, S., & Høiby, N. (2002). Inhibition of quorum sensing in *Pseudomonas aeruginosa* bioflm bacteria by a halogenated furanone compound. *Microbiology, 148*, 87–102.
- <span id="page-15-21"></span>136. Lönn-Stensrud, J., Petersen, F., Benneche, T., & Scheie, A. A. (2007). Synthetic bromated furanone inhibits autoinducer-2-mediated communication and bioflm formation in oral streptococci. *Oral Microbiology and Immunology, 22*, 340–346.
- <span id="page-15-22"></span>137. O'Loughlin, C. T., Miller, L. C., Siryaporn, A., Drescher, K., Semmelhack, M. F., & Bassler, B. L. (2013). A quorum-sensing inhibitor blocks *Pseudomonas aeruginosa* virulence and bioflm formation. *Proceedings of the National Academy of Sciences, 110*, 17981–17986.
- <span id="page-15-23"></span>138. Ishii, S., Fukui, K., Yokoshima, S., Kumagai, K., Beniyama, Y., Kodama, T., Fukuyama, T., Okabe, T., Nagano, T., & Kojima, H. (2017). High-throughput screening of small molecule inhibitors of the Streptococcus quorum-sensing signal pathway. *Science and Reports, 7*, 1–10.
- <span id="page-15-24"></span>139. Heidari, A., Noshiranzadeh, N., Haghi, F., & Bikas, R. (2017). Inhibition of quorum sensing related virulence factors of *Pseudomonas aeruginosa* by pyridoxal lactohydrazone. *Microbial Pathogenesis, 112*, 103–110.
- <span id="page-15-25"></span>140. Heidari, A., Haghi, F., Noshiranzadeh, N., & Bikas, R. (2017). (S, E)-2-hydroxy-N-(2-hydroxy-5-nitrobenzylidene) propane hydrazide as a quorum sensing inhibitor of *Pseudomonas aeruginosa*. *Medicinal Chemistry Research, 26*, 1947–1955.
- <span id="page-15-0"></span>141. Alizadeh, N., Memar, M., Mehramuz, B., Abibiglou, S., Hemmati, F., & Samadi Kafil, H. (2018). Current advances in aptamer-assisted technologies for detecting bacterial and fungal toxins. *Journal of Applied Microbiology, 124*, 644–651.
- <span id="page-15-1"></span>142. Ning, Y., Cheng, L., Ling, M., Feng, X., Chen, L., Wu, M., & Deng, L. (2015). Efficient suppression of biofilm formation by a nucleic acid aptamer. *Pathogens and Disease, 73*, ftv034.
- <span id="page-15-2"></span>143. Nimjee, S. M., Rusconi, C. P., & Sullenger, B. A. (2005). Aptamers: An emerging class of therapeutics. *Annual Review of Medicine, 56*, 555–583.
- <span id="page-15-3"></span>144. Shatila, F., Yaşa, İ, & Yalçın, H. T. (2020). Inhibition of *Salmonella enteritidis* bioflms by Salmonella invasion protein-targeting aptamer. *Biotechnology Letters*. [https://doi.org/10.1007/](https://doi.org/10.1007/s10529-020-02920-2) [s10529-020-02920-2](https://doi.org/10.1007/s10529-020-02920-2)
- <span id="page-15-4"></span>145. Kolovskaya, O. S., Savitskaya, A. G., Zamay, T. N., Reshetneva, I. T., Zamay, G. S., Erkaev, E. N., Wang, X., Wehbe, M., Salmina, A. B., & Perianova, O. V. (2013). Development of bacteriostatic DNA aptamers for salmonella. *Journal of Medicinal Chemistry, 56*, 1564–1572.
- <span id="page-15-5"></span>146. Mao, B., Cheng, L., Wang, S., Zhou, J., & Deng, L. (2018). Combat bioflm by bacteriostatic aptamer-functionalized graphene oxide. *Biotechnology and Applied Biochemistry, 65*, 355–361.
- <span id="page-15-6"></span>147. Lijuan, C., Xing, Y., Minxi, W., Wenkai, L., & Le, D. (2017). Development of an aptamer-ampicillin conjugate for treating bioflms. *Biochemical and Biophysical Research Communications, 483*, 847–854.
- <span id="page-15-7"></span>148. Thevendran, R., Sarah, S., Tang, T.-H., & Citartan, M. (2020). Strategies to bioengineer aptamer-driven nanovehicles as exceptional molecular tools for targeted therapeutics: A review. *Journal of Controlled Release*. [https://doi.org/10.1016/j.jconrel.2020.](https://doi.org/10.1016/j.jconrel.2020.04.051) [04.051](https://doi.org/10.1016/j.jconrel.2020.04.051)
- <span id="page-15-8"></span>149. Yu, Y. M., Xu, B. Y., Yan, S. S., Xu, J. F., Liu, F., Li, G. M., Ding, Y. L., & Wu, S. Q. (2013). Screening and anti-virulent study of N-acyl homoserine lactones DNA aptamers against *Pseudomonas aeruginosa* quorum sensing. *Biotechnology and Bioprocess Engineering, 18*, 406–412.
- <span id="page-15-9"></span>150. Wang, S., Mao, B., Wu, M., Liang, J., & Deng, L. (2018). Infuence of aptamer-targeted antibioflm agents for treatment of *Pseudomonas aeruginosa* bioflms. *Antonie van Leeuwenhoek, 111*, 199–208.
- <span id="page-15-26"></span>151. Oroh, S. B., Mustopa, A. Z., Budiarti, S., & Budiarto, B. R. (2020). Inhibition of enteropathogenic *Escherichia coli* bioflm formation by DNA aptamer. *Molecular Biology Reports*. [https://](https://doi.org/10.1007/s11033-020-05822-8) [doi.org/10.1007/s11033-020-05822-8](https://doi.org/10.1007/s11033-020-05822-8)
- <span id="page-15-27"></span>152. Sengupta, B., Adhikari, P., Mallet, E., Havner, R., & Pradhan, P. (2020). Spectroscopic study on *Pseudomonas aeruginosa* bioflm in the presence of the Aptamer-DNA scafolded silver nanoclusters. *Molecules, 25*, 3631.
- <span id="page-15-28"></span>153. Whitesides, G. M. (2005). Nanoscience, nanotechnology, and chemistry. *Small (Weinheim an der Bergstrasse, Germany), 1*, 172–179.
- <span id="page-15-29"></span>154. Chen, M., Yu, Q., & Sun, H. (2013). Novel strategies for the prevention and treatment of bioflm related infections. *International Journal of Molecular Sciences, 14*, 18488–18501.
- <span id="page-15-30"></span>155. Mahamuni-Badiger, P. P., Patil, P. M., Badiger, M. V., Patel, P. R., Thorat-Gadgil, B. S., Pandit, A., & Bohara, R. A. (2019). Bioflm formation to inhibition: Role of zinc oxide-based nanoparticles. *Materials Science and Engineering: C, 108*, 110319.
- <span id="page-16-0"></span>156. Shi, S.-F., Jia, J.-F., Guo, X.-K., Zhao, Y.-P., Chen, D.-S., Guo, Y.-Y., & Zhang, X.-L. (2016). Reduced *Staphylococcus aureus* bioflm formation in the presence of chitosan-coated iron oxide nanoparticles. *International JOURNAL of Nanomedicine, 11*, 6499.
- <span id="page-16-1"></span>157. Gupta, D., Singh, A., & Khan, A. U. (2017). Nanoparticles as efflux pump and biofilm inhibitor to rejuvenate bactericidal effect of conventional antibiotics. *Nanoscale Research Letters, 12*, 1–6.
- <span id="page-16-2"></span>158. Padwal, P., Bandyopadhyaya, R., & Mehra, S. (2014). Polyacrylic acid-coated iron oxide nanoparticles for targeting drug resistance in mycobacteria. *Langmuir, 30*, 15266–15276.
- <span id="page-16-3"></span>159. Banoee, M., Seif, S., Nazari, Z. E., Jafari-Fesharaki, P., Shahverdi, H. R., Moballegh, A., Moghaddam, K. M., & Shahverdi, A. R. (2010). ZnO nanoparticles enhanced antibacterial activity of ciprofoxacin against *Staphylococcus aureus* and *Escherichia coli*. *Journal of Biomedical Materials Research Part B: Applied Biomaterials, 93*, 557–561.
- <span id="page-16-4"></span>160. Nallathamby, P. D., Lee, K. J., Desai, T., & Xu, X.-H.N. (2010). Study of the multidrug membrane transporter of single living *Pseudomonas aeruginosa* cells using size-dependent plasmonic nanoparticle optical probes. *Biochemistry, 49*, 5942–5953.
- <span id="page-16-5"></span>161. Pérez-Laguna, V., García-Luque, I., Ballesta, S., Pérez-Artiaga, L., Lampaya-Pérez, V., Samper, S., Soria-Lozano, P., Rezusta, A., & Gilaberte, Y. (2018). Antimicrobial photodynamic activity of Rose Bengal, alone or in combination with Gentamicin, against planktonic and bioflm *Staphylococcus aureus*. *Photodiagnosis and Photodynamic Therapy, 21*, 211–216.
- <span id="page-16-6"></span>162. Khan, F., Lee, J.-W., Manivasagan, P., Pham, D. T. N., Oh, J., & Kim, Y.-M. (2019). Synthesis and characterization of chitosan oligosaccharide-capped gold nanoparticles as an efective antibioflm drug against the *Pseudomonas aeruginosa* PAO1. *Microbial Pathogenesis, 135*, 103623.
- <span id="page-16-7"></span>163. Wang, Z., Bai, H., Lu, C., Hou, C., Qiu, Y., Zhang, P., Duan, J., & Mu, H. (2019). Light controllable chitosan micelles with ROS generation and essential oil release for the treatment of bacterial bioflm. *Carbohydrate Polymers, 205*, 533–539.
- <span id="page-16-8"></span>164. Kavanaugh, J. S., Flack, C. E., Lister, J., Ricker, E. B., Ibberson, C. B., Jenul, C., Moormeier, D. E., Delmain, E. A., Bayles, K. W., & Horswill, A. R. (2019). Identifcation of extracellular DNA-binding proteins in the bioflm matrix. *MBio, 10*, e01137–e1119.
- 165. Xia, T., Kovochich, M., Liong, M., Madler, L., Gilbert, B., Shi, H., Yeh, J. I., Zink, J. I., & Nel, A. E. (2008). Comparison of the mechanism of toxicity of zinc oxide and cerium oxide nanoparticles based on dissolution and oxidative stress properties. *ACS Nano, 2*, 2121–2134.
- 166. Zhang, L., Jiang, Y., Ding, Y., Povey, M., & York, D. (2007). Investigation into the antibacterial behaviour of suspensions of ZnO nanoparticles (ZnO nanofuids). *Journal of Nanoparticle Research, 9*, 479–489.
- <span id="page-16-9"></span>167. Ebrahimzadeh, S., Bari, M. R., Hamishehkar, H., Kafl, H. S., & Lim, L.-T. (2021). Essential oils-loaded electrospun chitosanpoly(vinyl alcohol) nonwovens laminated on chitosan flm as bilayer bioactive edible flms. *LWT, 144*, 111217.
- <span id="page-16-10"></span>168. Ma, L., Conover, M., Lu, H., Parsek, M. R., Bayles, K., & Wozniak, D. J. (2009). Assembly and development of the *Pseudomonas aeruginosa* bioflm matrix. *PLoS Pathogens, 5*, e1000354.
- <span id="page-16-11"></span>169. Whitchurch, C. B., Tolker-Nielsen, T., Ragas, P. C., & Mattick, J. S. (2002). Extracellular DNA required for bacterial bioflm formation. *Science, 295*, 1487–1487.
- <span id="page-16-12"></span>170. Kovach, K., Fleming, D., Rumbaugh, K. P., & Gordon, V. (2019). Specifc disruption of established *P. aeruginosa* bioflms using polymer-attacking enzymes. *bioRxiv*, 598979.
- <span id="page-16-13"></span>171. Zhang, L., Jiang, Y., Ding, Y., Daskalakis, N., Jeuken, L., Povey, M., O'Neill, A. J., & York, D. W. (2010). Mechanistic

investigation into antibacterial behaviour of suspensions of ZnO nanoparticles against *E. coli*. *Journal of Nanoparticle Research, 12*, 1625–1636.

- <span id="page-16-14"></span>172. Messiaen, A.-S., Forier, K., Nelis, H., Braeckmans, K., & Coenye, T. (2013). Transport of nanoparticles and tobramycinloaded liposomes in *Burkholderia cepacia* complex bioflms. *PLoS ONE, 8*, e79220.
- <span id="page-16-17"></span>173. Pham, D. T. N., Khan, F., Phan, T. T. V., Park, S.-K., Manivasagan, P., Oh, J., & Kim, Y.-M. (2019). Biofilm inhibition, modulation of virulence and motility properties by FeOOH nanoparticle in *Pseudomonas aeruginosa*. *Brazilian Journal of Microbiology, 50*, 791–805.
- <span id="page-16-18"></span>174. Loo, C. Y., Rohanizadeh, R., Young, P. M., Traini, D., Cavaliere, R., Whitchurch, C. B., & Lee, W. H. (2016). Combination of silver nanoparticles and curcumin nanoparticles for enhanced anti-bioflm activities. *Journal of Agriculture and Food Chemistry, 64*, 2513–2522.
- <span id="page-16-19"></span>175. Ghotaslou, R., Bahari, Z., Aliloo, A., Gholizadeh, P., & Eshlaghi, B. S. (2017). The in vitro efects of silver nanoparticles on bacterial bioflms. *Journal of Microbiology, Biotechnology and Food Sciences, 6*, 1077–1080.
- <span id="page-16-20"></span>176. Mu, H., Tang, J., Liu, Q., Sun, C., Wang, T., & Duan, J. (2016). Potent antibacterial nanoparticles against bioflm and intracellular bacteria. *Science and Reports, 6*, 1–9.
- <span id="page-16-21"></span>177. Lin, W.-T., Tan, H.-L., Duan, Z.-L., Yue, B., Ma, R., He, G., & Tang, T.-T. (2014). Inhibited bacterial bioflm formation and improved osteogenic activity on gentamicin-loaded titania nanotubes with various diameters. *International Journal of Nanomedicine, 9*, 1215.
- <span id="page-16-22"></span>178. Khan, F., Manivasagan, P., Lee, J.-W., Pham, D. T. N., Oh, J., & Kim, Y.-M. (2019). Fucoidan-stabilized gold nanoparticle-mediated bioflm inhibition, attenuation of virulence and motility properties in *Pseudomonas aeruginosa* PAO1. *Marine Drugs, 17*, 208.
- <span id="page-16-23"></span>179. Ali, S. G., Ansari, M. A., Khan, H. M., Jalal, M., Mahdi, A. A., & Cameotra, S. S. (2017). *Crataeva nurvala* nanoparticles inhibit virulence factors and bioflm formation in clinical isolates of *Pseudomonas aeruginosa*. *Journal of Basic Microbiology, 57*, 193–203.
- <span id="page-16-24"></span>180. Huang, J., Liu, Y., Yang, L., & Zhou, F. (2019). Synthesis of sulfonated chitosan and its antibioflm formation activity against *E. coli* and *S. aureus*. *International Journal of Biological Macromolecules, 129*, 980–988.
- <span id="page-16-25"></span>181. Khan, S. T., Ahmad, J., Ahamed, M., Musarrat, J., & Al-Khedhairy, A. A. (2016). Zinc oxide and titanium dioxide nanoparticles induce oxidative stress, inhibit growth, and attenuate bioflm formation activity of Streptococcus mitis. *JBIC Journal of Biological Inorganic Chemistry, 21*, 295–303.
- <span id="page-16-15"></span>182. Li, X., Wong, C.-H., Ng, T.-W., Zhang, C.-F., Leung, K.C.- F., & Jin, L. (2016). The spherical nanoparticle-encapsulated chlorhexidine enhances anti-biofilm efficiency through an efective releasing mode and close microbial interactions. *International Journal of Nanomedicine, 11*, 2471.
- <span id="page-16-16"></span>183. Wojciechowska, M., Równicki, M., Mieczkowski, A., Miszkiewicz, J., & Trylska, J. (2020). Antibacterial peptide nucleic acids—Facts and perspectives. *Molecules, 25*, 559.
- <span id="page-16-26"></span>184. Lee, H. T., Kim, S. K., & Yoon, J. W. (2019). Antisense peptide nucleic acids as a potential anti-infective agent. *Journal of Microbiology, 57*, 423–430.
- <span id="page-16-27"></span>185. Kurupati, P., Tan, K. S. W., Kumarasinghe, G., & Poh, C. L. (2007). Inhibition of gene expression and growth by antisense peptide nucleic acids in a multiresistant β-lactamase-producing *Klebsiella pneumoniae* strain. *Antimicrobial Agents and Chemotherapy, 51*, 805–811.
- <span id="page-16-28"></span>186. Barkowsky, G., Lemster, A.-L., Pappesch, R., Jacob, A., Krüger, S., Schröder, A., Kreikemeyer, B., & Patenge, N.

(2019). Infuence of diferent cell-penetrating peptides on the antimicrobial efficiency of PNAs in *Streptococcus pyogenes*. *Molecular Therapy-Nucleic Acids, 18*, 444–454.

- <span id="page-17-0"></span>187. Narenji, H., Gholizadeh, P., Aghazadeh, M., Rezaee, M. A., Asgharzadeh, M., & Kafl, H. S. (2017). Peptide nucleic acids (PNAs): Currently potential bactericidal agents. *Biomedicine & Pharmacotherapy, 93*, 580–588.
- <span id="page-17-1"></span>188. Readman, J. B., Dickson, G., & Coldham, N. G. (2017). Tetrahedral DNA nanoparticle vector for intracellular delivery of targeted peptide nucleic acid antisense agents to restore antibiotic sensitivity in cefotaxime-resistant *Escherichia coli*. *Nucleic Acid Therapeutics, 27*, 176–181.
- <span id="page-17-2"></span>189. Otsuka, T., Brauer, A. L., Kirkham, C., Sully, E. K., Pettigrew, M. M., Kong, Y., Geller, B. L., & Murphy, T. F. (2016). Antimicrobial activity of antisense peptide–peptide nucleic acid conjugates against non-typeable *Haemophilus infuenzae* in planktonic and bioflm forms. *Journal of Antimicrobial Chemotherapy, 72*, 137–144.
- <span id="page-17-3"></span>190. Doyle, T. B., Hawkins, A. C., & McCarter, L. L. (2004). The complex fagellar torque generator of *Pseudomonas aeruginosa*. *Journal of Bacteriology, 186*, 6341–6350.
- <span id="page-17-4"></span>191. Xia, Y., Xiong, Y., Li, X., & Su, X. (2011). Inhibition of bioflm formation by the antisense peptide nucleic acids targeted at the motA gene in *Pseudomonas aeruginosa* PAO1 strain. *World Journal of Microbiology & Biotechnology, 27*, 1981–1987.
- <span id="page-17-5"></span>192. Castillo, J. I., Równicki, M., Wojciechowska, M., & Trylska, J. (2018). Antimicrobial synergy between mRNA targeted peptide nucleic acid and antibiotics in *E. coli*. *Bioorganic & Medicinal Chemistry Letters, 28*, 3094–3098.

**Publisher's Note** Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional afliations.