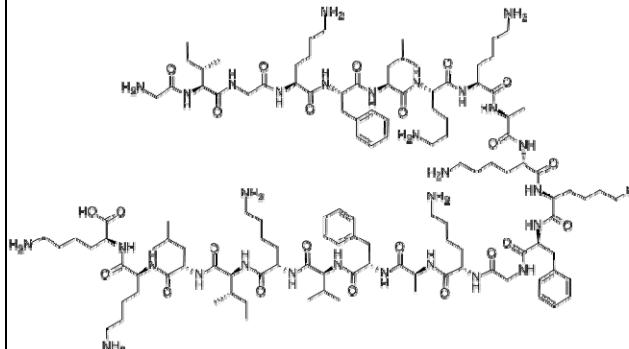
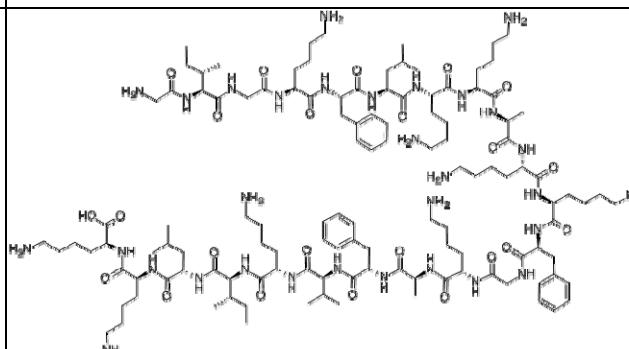
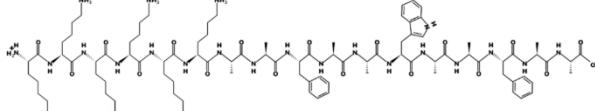
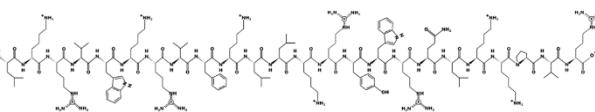
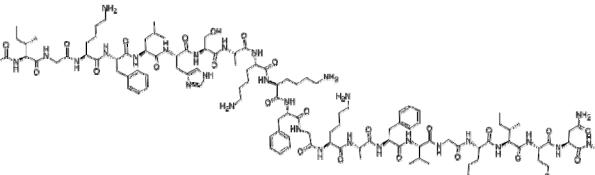
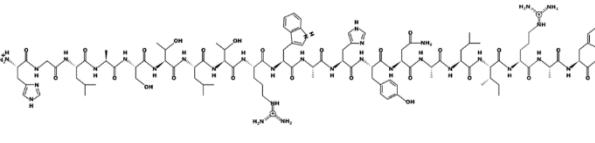
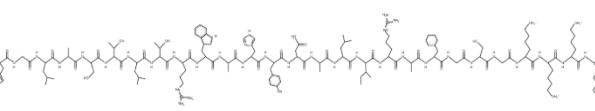
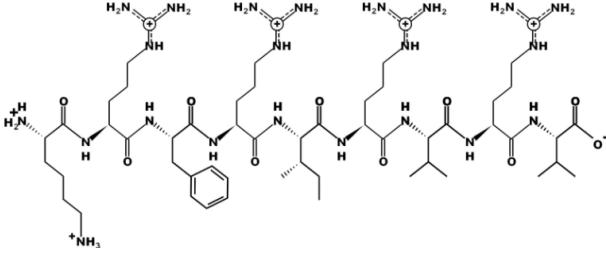
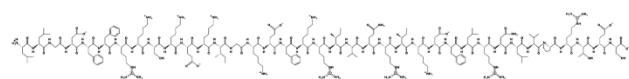
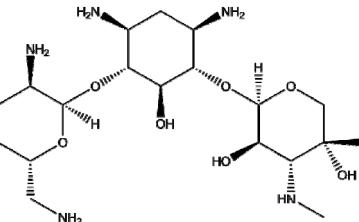
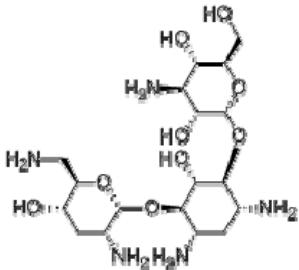


Approach	Molecular structure	Biofilm-forming pathogens	Source	Mechanism(s)	Dose/Effect	Reference(s)
Antimicrobial peptides		<i>P. aeruginosa</i> Z25.1	DFU	n.d.	MBIC = 80 µg/ml MBEC = 100 µg/ml	[182]
		<i>S. aureus</i> Z25.2	DFU	n.d.	MBIC = 10 µg/ml MBEC = 9 µg/ml	[182]
		<i>P. aeruginosa</i> + <i>S. aureus</i>	DFU	n.d.	MBIC = 50 µg/ml MBEC = 120 µg/ml	[182]
		<i>P. aeruginosa</i> Z25.1	DFU	n.d.	MBIC = 70 µg/ml MBEC = 90 µg/ml	[182]
		<i>S. aureus</i> Z25.2	DFU	n.d.	MBIC = 1 µg/ml MBEC = 5 µg/ml	[182]
		<i>P. aeruginosa</i> + <i>S. aureus</i>	DFU	n.d.	MBIC = 50 µg/ml MBEC = 50 µg/ml	[182]

	6K-F17		<i>P. aeruginosa</i> PAO1	Reference strain	Interaction with biofilm EPS	10 µg/ml (75% biofilm reduction)	[184]
SAAP-148		<i>A. baumannii</i>	Clinical isolates	Interaction and subsequently permeabilization of the cytoplasmic membrane, leading to bacterial death	12.8 mM (85% biofilm reduction)	[177]	
		<i>S. aureus</i>	Clinical isolates	12.8 mM (81% biofilm reduction)			
Magainin 2		<i>A. baumannii</i>	Clinical isolates	n.d.	256 µM (66% biofilm inhibition)	[178]	
gH625		<i>S. aureus</i>	Otitis infection	Biofilm penetration and interaction with microbial membranes. The modification of the gH625 peptide, which consists of the addition of a sequence of lysine residues, increased the capacity of penetration and interaction	50 µM (35% biofilm prevention)	[179]	
		Polimicrobial biofilms ( <i>Candida tropicalis</i> and <i>S. aureus</i> )	Systemic infection and otitis infection	50 µM (48% biofilm prevention)			
		<i>S. aureus</i>	Otitis infection	50 µM (50% biofilm prevention)			
		Polimicrobial biofilms ( <i>C. tropicalis</i> and <i>S. aureus</i> )	Systemic infection and otitis infection	50 µM (88% biofilm prevention)			

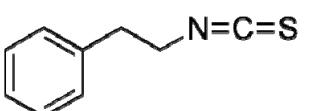
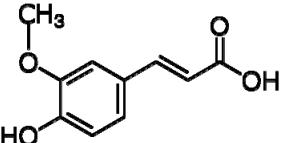
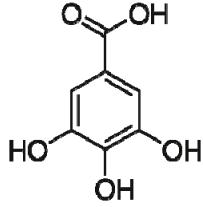
	Hp1404-T1e		<i>P. aeruginosa</i>	ATCC 27853	n.d.	MBIC = 12,5 µM	[180]
DJK-5		<i>P. aeruginosa</i>	Cystic fibrosis	Clinical isolates	Binding and promotion of degradation of the signal for biofilm formation and maintenance (p)ppGpp	MBIC <sub>50</sub> = 1 µg/ml	[192]
			<i>K. pneumoniae</i>			MBIC <sub>50</sub> : 8 to >64 µg/ml	[194]
		<i>K. pneumoniae</i>	Cystic fibrosis			MBIC <sub>50</sub> : 1 to 8 µg/ml	
DJK-6		<i>P. aeruginosa</i>	Cystic fibrosis	Clinical isolates	Binding and promotion of degradation of the signal for biofilm formation and maintenance (p)ppGpp	MBIC <sub>50</sub> =0,5µg/m l	[192]
			<i>S. epidermidis</i>			40 % biofilm reduction	
		<i>S. aureus</i>	-			49% biofilm reduction	[173]
FI-P <sub>R</sub> P <sub>R</sub> P <sub>L</sub> -5		<i>A. baumannii</i>	Clinical isolates	Binding and promotion of degradation of the signal for biofilm formation and maintenance (p)ppGpp	MBIC <sub>50</sub> = 2 µg/ml	[193]	
					MBIC <sub>50</sub> = 8 µg/ml		
		<i>E. coli</i>			MBIC <sub>50</sub> = 2 µg/ml		
		<i>K. pneumoniae</i>	ATTC 13883		MBIC <sub>50</sub> = 2 µg/ml		
		MRSA	Clinical isolates		MBIC <sub>50</sub> = 2 µg/ml		
		<i>P. aeruginosa</i> PAO1	Wild type		MBIC <sub>50</sub> = 5 µg/ml		

	1037		<i>P. aeruginosa</i> PAO1	Reference strain	Inhibition of swimming and swarming motilities and stimulation of twitching motility	10 µg/ml (50% biofilm inhibition)	[186]
	LL-37		<i>S. aureus</i> ATCC 25923	Affect the expression of <i>agr</i> locus and <i>ica</i> locus genes, essential for the development of biofilm	10 µg/ml (40% biofilm inhibition)	[172]	
			<i>P. aeruginosa</i> PAO1	Reference strain	Decrease attachment, increase twitching motility and affect <i>Las</i> and <i>Rhl</i> QS systems	16 µg/ml (75% biofilm reduction)	[185]
Bacteriophages	vB_PaeM_LS1		<i>P. aeruginosa</i>	Human isolates	98% biofilm reduction (from 7.3 to 4.8 log <sub>10</sub> CFU/ml)	[200]	
	ZCKP1		<i>K. pneumoniae</i>	DFU	Biofilm bacteria are susceptible to lytic bacteriophages  ≈ 60 – 90% biofilm viability after 24h, single treatment	[204]	
	NP1 + NP3 + Gentamicin		<i>P. aeruginosa</i>	-	1.5-log reduction	[208]	

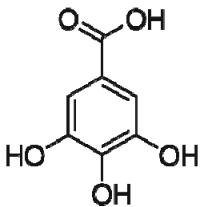
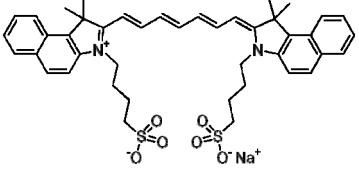
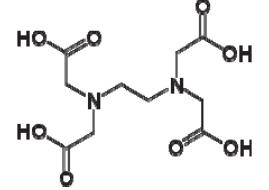
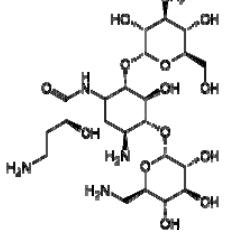
	NP1 + NP3 + Tobramycin		<i>P. aeruginosa</i>	-	2-log reduction	[208]
	vB_AbaM-IME-AB2		<i>A. baumannii</i>	Expectoration	88.5% biofilm inhibition ( $5.6 \times 10^{-7}$ CFU)	[199]
	Bacteriophage cocktail (DL52, DL54, DL60, DL62, DL64 and DL68)		<i>P. aeruginosa</i> PAO1	Reference strain	95% biofilm reduction after 4h	[206]
	DRA88 + phage K		<i>S. aureus</i>	Clinical isolates	100% biofilm disruption after 48h	[197]
	Bacteriophage cocktail ( <i>S. aureus</i> F44/10 and F125/10; <i>P. aeruginosa</i> F770/05 and F510/08; <i>A. baumannii</i> F1245/05)		<i>A. baumannii</i>	Clinical isolates	60% biofilm reduction after 24h, one-time bacteriophage application	[198]
			<i>P. aeruginosa</i>	Clinical isolates	50% biofilm reduction after 24h, one-time bacteriophage application	
			<i>S. aureus</i>	Clinical isolates	50% biofilm reduction after 24h, one-time bacteriophage application	
	AB7-IBB2		<i>A. baumannii</i>	Clinical isolates	80% biofilm reduction	[196]
	SAP-26		<i>S. aureus</i>	Clinical isolates	2-log reduction	[207]

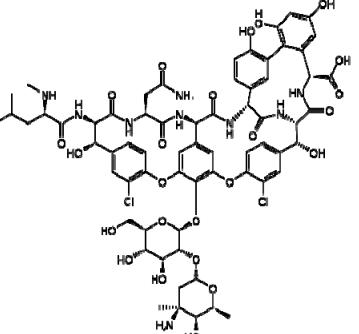
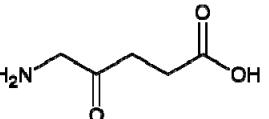
	SAP-26 + Rifampicin		<i>S. aureus</i>	Clinical isolates		4-log reduction	[207]
Phytochemicals	$\alpha$ -mangostin		<i>P. aeruginosa</i>	DFU	n.d.	2 $\mu\text{g/ml}$	[169]
			<i>S. epidermidis</i>	DFU	n.d.	2 $\mu\text{g/ml}$	[169]
	Lutein		<i>P. aeruginosa</i> PAO1	Reference strain	n.d.	20 $\mu\text{g/ml}$ (61% biofilm inhibition)	[167]
	Quercetin		<i>P. aeruginosa</i> PAO1	Reference strain	Transcriptional changes associated with QS and expression levels of lasI, lasR, rhlI and rhlR significantly reduced	250 $\mu\text{g/ml}$ (43 – 78% biofilm inhibition)	[213]
	Caffeic acid		MRSA	clinical isolates	Interference with bacterial motility, changes in the physico-chemical properties of cell surfaces which alter the adhesion potential to PS, and interference with QS; and hinder the biofilm formation process	16 $\mu\text{g/ml}$ (51% biofilm inhibition)	[219]
	<i>p</i> -coumaric acid					24% biofilm reduction	[212]
	Trans-cinnamaldehyde					28% biofilm reduction	[212]
						42% biofilm reduction; 79% inactivation	[212]

	7-hydroxycoumarin		<i>E. coli</i>	CETC 434		800 µg/ml (51% biofilm reduction; 72% biofilm inactivation)	[217]
			<i>S. aureus</i>	CETC 976		200 µg/ml (10% biofilm reduction; 40% biofilm inactivation)	[217]
	Indole-3-carbinol		<i>E. coli</i>	CETC 434		800 µg/ml (43% biofilm reduction; 75% biofilm inactivation)	[217]
			<i>S. aureus</i>	CETC 976		400 µg/ml (47% biofilm inactivation)	[217]
	Salicylic acid		<i>E. coli</i>	CETC 434		3200 µg/ml (34% biofilm reduction; 75% biofilm inactivation)	[217]
			<i>S. aureus</i>	CETC 976		1600 µg/ml (47% biofilm inactivation)	[217]
	Saponin		<i>E. coli</i>	CETC 434		3200 µg/ml (34% biofilm reduction; 75% biofilm inactivation)	[217]
			<i>S. aureus</i>	CETC 976		3200 µg/ml (48% biofilm inactivation)	[217]
	Allyl isothiocyanate		<i>E. coli</i>	CETC 434		1000 µg/ml (87% biofilm prevention)	[210]
			<i>P. aeruginosa</i>	ATCC 10145		1000 µg/ml (99% biofilm prevention)	[210]
			<i>S. aureus</i>	CETC 976		1000 µg/ml (96% biofilm prevention)	[210]

	2-phenylethyl isothiocyanate		<i>E. coli</i>	CETC 434		1000 µg/ml (100% biofilm prevention)	[210]
			<i>P. aeruginosa</i>	ATCC 10145		1000 µg/ml (93% biofilm prevention)	[210]
			<i>S. aureus</i>	CETC 976		1000 µg/ml (90% biofilm prevention)	[210]
			MRSA	Clinical isolates		27% biofilm reduction	[212]
	Ferulic acid		<i>E. coli</i>	CETC 434		2500 µg/ml (13% biofilm prevention; 38% biofilm reduction)	[216]
			<i>P. aeruginosa</i>	ATCC 10145		500 µg/ml (81% biofilm prevention; 42% biofilm reduction)	[216]
			<i>S. aureus</i>	CETC 976		5000 µg/ml (6% biofilm prevention)	[216]
	Gallic acid		<i>E. coli</i>	CETC 434		5000 µg/ml (64% biofilm prevention; 29% biofilm reduction)	[216]
			<i>P. aeruginosa</i>	ATCC 10145		500 µg/ml (84% biofilm prevention; 48% biofilm reduction)	[216]
			<i>S. aureus</i>	CETC 976		500 µg/ml (70% biofilm prevention)	[216]
Phytosynthesized silver nanoparticles (AgNPs)	<i>Bridelia retusa</i> mediated AgNPs (BR-AgNPs)		<i>E. coli</i>	MTCC 739	Interference with QS	MBIC <sub>50</sub> = 64 µg/ml	[224]
			<i>P. aeruginosa</i>	MTCC 741		MBIC <sub>50</sub> = 23 µg/ml	[224]
			<i>S. aureus</i>	MTCC 96		MBIC <sub>50</sub> = 33 µg/ml	[224]

	<i>Glochidion lanceolarium</i> mediated AgNPs (GL-AgNPs)		<i>E. coli</i>	MTCC 739		MBIC <sub>50</sub> = 46 µg/ml	[224]
			<i>P. aeruginosa</i>	MTCC 741		MBIC <sub>50</sub> = 69 µg/ml	[224]
			<i>S. aureus</i>	MTCC 96		MBIC <sub>50</sub> = 53 µg/ml	[224]
	<i>Semecarpus anacardium</i> mediated AgNPs (SA-AgNPs)		<i>E. coli</i>	MTCC 739		MBIC <sub>50</sub> = 23 µg/ml	[224]
			<i>P. aeruginosa</i>	MTCC 741		MBIC <sub>50</sub> = 13 µg/ml	[224]
			<i>S. aureus</i>	MTCC 96		MBIC <sub>50</sub> = 34 µg/ml	[224]
	<i>Aerva lanata</i> mediated AgNPs (AL-AgNPs)		<i>B. subtilis</i>			MIC (10 µg/ml; 35% biofilm disruption) MBC (15 µg/ml; 89% biofilm disruption)	[229]
			<i>E. coli</i>			MIC (5 µg/ml; 79% biofilm disruption) MBC (10 µg/ml; 99% biofilm disruption)	[229]
			<i>P. aeruginosa</i>			MIC (10 µg/ml; 45% biofilm disruption) MBC (15 µg/ml; 82% biofilm disruption)	[229]
			<i>S. aureus</i>			MIC (15 µg/ml; 47% biofilm disruption) MBC (20 µg/ml; 86% biofilm disruption)	[229]
	<i>Piper betle</i> mediated AgNPs (Pb-AgNPs)		<i>P. aeruginosa</i> PAO1	Reference strain	Inactivation of transcriptional regulator <i>lasR</i> and inhibition of AHL production	8 µg/ml (78 % biofilm inhibition)	[223]

	Gallic acid mediated AgNPs (GA-AgNPs)		<i>P. aeruginosa</i> PAO1	Reference strain	n.d.	biofilm under high fluid shears conditions (100 mg/ml; ≈ 4-log reduction)	[228]
						biofilm under static conditions (500 mg/ml; ≈ 4-log reduction)	[228]
Photodynamic therapy	PDT-ICG		MDR <i>P. aeruginosa</i>	Clinical isolates	PDT generate a large amount of ROS, which then acted on bacterial cell walls, membranes and nucleic acids	MBIC (16 µg/ml; 90% survival ratio)	[36]
			MRSA			MBIC (32 µg/ml; 61% survival ratio)	[36]
	PDT-ICG + EDTA		MDR <i>P. aeruginosa</i>			MBIC (16 µg/ml; 53% survival ratio)	[36]
			MRSA			MBIC (32 µg/ml; 55% survival ratio)	[36]
	PDT-ICG + EDTA		MDR <i>P. aeruginosa</i>		EDTA counteracts biofilms by chelating Mg²⁺ and Ca²⁺; remove iron atoms that are essential for virulence and pathogenicity	MBIC (16 µg/ml; 12% survival ratio)	[36]

		Vancomycin		MRSA		MBIC (32 µg/ml; 13% survival ratio)	[36]	
PDT-5-ALA			<i>S. aureus</i>	Clinical isolates PDT generate a large amount of ROS, which then acted on bacterial cell walls, membranes and nucleic acids		500 J/cm <sup>2</sup> (20% survival ratio)	[234]	
PDT-5-ALA			<i>S. epidermidis</i>			500 J/cm <sup>2</sup> (20% survival ratio)	[234]	
PDT-5-ALA			<i>S. aureus</i>			500 J/cm <sup>2</sup> + 2 µg/ml (~15% survival ratio)	[234]	
Gentamicin			<i>S. epidermidis</i>			500 J/cm <sup>2</sup> + 2 µg/ml (~10% survival ratio)	[234]	

	PDT-TMP	Vancomycin	<p>The image contains two chemical structures. The top structure is PDT-TMP, which is a porphyrin derivative consisting of four pyrrole rings linked by methine groups, with various substituents including phenyl groups and methyl groups. The bottom structure is Vancomycin, a complex polypeptide antibiotic composed of two peptide chains linked by a disulfide bond, featuring multiple amide linkages and hydroxyl groups.</p>	MRSA	Clinical isolates	EPS disruption	150 J/cm <sup>2</sup> + 10 µM TMP + 1 µg/ml	[235]
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Minimal biofilm inhibitory concentration (MBIC); Minimal biofilm eradication concentration (MBEC); Minimal Biofilm Inhibitory Concentration leading to 50% decrease in biofilm growth (MBIC50); n.d. – not determined; American Type Culture Collection (ATCC); Spanish Type Culture Collection (CETC); Microbial Type Culture Collection (MTCC); Tetramesitylporphyrin (TMP)