### Improving phage therapy by evasion of phage resistance mechanisms

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Antibiotic failure is one of the most worrisome threats to global health. Among the new therapeutic efforts that are being explored, the use of bacteriophages (viruses that kill bacteria), also known as 'phages', is being extensively studied as a strategy to target bacterial pathogens. However, one of the main drawbacks of phage therapy is the plethora of defence mechanisms that bacteria use to defend themselves against phages. This review aims to summarize the therapeutic approaches that are being evaluated to overcome the bacterial defence systems, including the most innovative therapeutic approaches applied: circumvention of phage receptor mutations; modification of prophages; targeting of CRISPR-Cas systems and the biofilm matrix; engineering of safer and more efficacious phages; and inhibition of the anti-persister strategies used by bacteria.

#### Introduction

Antimicrobial resistance causes almost 5 million deaths annually worldwide and is predicted to become the leading cause of death. Bacterial antibiotic resistance is driven by natural evolution, and antibiotic resistance genes are common, even in DNA isolated from ancient sediments. However, the massive use of antibacterial agents over decades, together with their release in untreated wastewater, exerts a selective pressure that has led to a global health crisis and could lead to an era without effective antibiotics. 4,5

Throughout history, numerous different outbreaks of severe infectious diseases have been caused by heterogeneous pathogens (belonging to different phyla) that have acquired some resistance mechanisms. Examples of such bacterial pathogens include carbapenem-resistant *Klebsiella pneumoniae*, <sup>6,7</sup> colistin-resistant plasmid-mediated Enterobacteriaceae, <sup>8–10</sup> XDR *Pseudomonas aeruginosa*, <sup>11</sup> carbapenem-resistant *Acinetobacter baumannii*, <sup>12</sup> MDR *Stenotrophomonas maltophilia*, <sup>13–15</sup> MDR *Burkholderia cepacia* <sup>13</sup> and MRSA. <sup>16,17</sup> The latter are some of the most common examples, but there are many more.

Although some researchers remain optimistic about a renewed antibiotic pipeline, vaccines or antibody–antibiotic conjugates, <sup>18</sup> every time a new antibiotic is introduced in the clinical

setting, bacterial resistance arises against it as the result of accelerated evolution.<sup>19</sup>

One of the most recently discovered antibiotics, teixobactin, was detected in 2015 by screening uncultured soil bacteria by using iChip technology.<sup>20</sup>

Teixobactin is a depsipeptide synthesized by the soil bacterium *Eleftheria terrae* and has shown extremely good efficacy and toxicity profiles *in vivo* against Gram-positive bacterial infections.<sup>21</sup>

To date, the resistance reported against teixobactin is slow and very costly in terms of fitness;<sup>22</sup> however, such successes are rare. The restoration of old strategies to combat resistant pathogens (such as phage therapy), together with the use of non-antibiotic compounds [antimicrobial peptides (AMPs) and repurposed drugs], is gaining much attention.<sup>23,24</sup>

#### **Bacteriophages**

Bacteriophages, also called phages, are viruses that infect bacteria. They are the most abundant organisms on Earth, being found in all environments including the soil, ocean, sewage and mammalian gut. It is thought that there are around 10 phages per bacterium, yielding an estimated 10<sup>31</sup> viral particles.<sup>25</sup> Phages are generally classified according to their life cycle into

lytic phages and lysogenic phages, although other types of phage infection are also possible, such as chronic infection or pseudolysogeny. <sup>26,27</sup> Lytic phages attach to the bacterial surface, inject their DNA, use bacterial machinery to replicate and subsequently lyse the cells to release multiple viral particles; by contrast, lysogenic phages inject their DNA and integrate it into the bacterial genome, so that it is passed on to progeny, where it may act as a vector for horizontal gene transfer (HGT). However, lysogenic phages can be induced and released from cells under stress situations, provoking cell lysis. <sup>28</sup>

#### Bacteria-phage interactions: an arms race

Bacteria and phages are in a permanent state of co-evolution referred to as an arms race, because when one develops a mechanism to evade the other, this causes the latter to adapt and avoid this defence system (Figure 1). Thus, bacteria have evolved defence mechanisms to protect themselves from predators, while phages, in turn, have evolved counterdefence strategies to evade these systems. <sup>29–32</sup> The main mechanisms of defence that bacteria have developed against phages are summarized in Table 1, and the phage resistance mechanisms are summarized in Table 2.

The use of phages for the apeutic purposes is of renewed interest in Western countries, where it fell into disuse in the 1940s with the advent of commercial antibiotics. Due to the rising antibiotic resistance rates, the use of phages in therapy has regained interest. Currently, the following minimum criteria are reguired for the use of phages for therapeutic purposes: (i) the phages must be strictly lytic in nature, lacking genes for lysogeny such as integrases and recombinases; (ii) they must have clear antimicrobial activity against the target pathogen; and (iii) bacterial debris and endotoxins must be removed to below predetermined thresholds of safety.<sup>56</sup> Although common in some eastern countries such as Georgia (Eastern Europe), phage therapy was completely abandoned with the advent of antibiotics in Western countries.<sup>57</sup> However, phage therapy has shown promising results, good safety profiles and efficacy in some relevant clinical cases, 58-61 as discussed below. Importantly, a recent retrospective and observational study focused on 100 consecutive cases of personalized phage therapy in Belgium revealed that more than 77% of difficult-to-treat infections experienced some clinical improvement, with bacterial eradication achieved in 61% of the total cases considered, and only 15% presenting adverse events (among which, 7% were considered mild to moderate).<sup>62</sup> Similarly, in 2023, Green et al.<sup>63</sup> summarized several cases of patients treated with personalized phage therapy and also concluded that there were no major adverse reactions and phage-antibiotic synergy (PAS) was frequently observed, consistent with the previously mentioned 2023 report by Pirnay et al.<sup>62</sup>

PAS is a well-known phenomenon consisting of an improved outcome after the combination of phage and antibiotics, relative to their separate effects. <sup>64,65</sup>

The synergism between conventional antibiotics and a preadapted lytic phage (*K. pneumoniae* phage M1) was made use of to resolve a severe fracture-related infection in a 30-year-old bomb victim in Belgium.<sup>65</sup> Pre-adapted phages were considered those mutants with broadest infectivity after 15 rounds of the Appelman protocol, which consists of growing phages iteratively



**Figure 1.** Original digital illustration (made in Adobe Photoshop from a sketch) representing the bacteria-phage 'arms race', a co-evolutionary state in which one develops a mechanism to evade the other and adapt. Bacteria display defence mechanisms to protect themselves from phages, while these evolve counterdefence strategies to evade these systems.

on mostly refractory bacterial isolates, until the adapted phage can lyse the phage-resistant strains. 66 Although ceftazidime/avibactam reduced K. pneumoniae bacterial counts in mature biofilms, they did not completely eradicate them, and high doses of the lytic phage M1 alone failed. Nonetheless, combinations of phage M1 and moderate concentrations of ceftazidime/avibactam were significantly more effective, which also suggests a synergistic effect. 65 Similarly, P. aeruginosa phage PNM and colistin (0.5 ma/L), aztreonam (8 ma/L) and gentamicin (2 ma/L) displayed strong synergistic activity against one of the P. aeruginosa isolates that caused severe post-liver transplantation sepsis in a toddler.<sup>67</sup> A similar synergistic phenomenon was observed with another P. aeruginosa isolate, although the effect was slightly less intense with colistin. Finally, the treatments allowed a second liver transplantation and resulted in complete resolution of the infection.<sup>67</sup> The most significant case reports using phages against antibiotic-resistant isolates are summarized in Table 3.

The strategies of phage training (pre-adaptation) and PAS do not always produce the intended results, which is why it is necessary to look for methods that specifically circumvent bacterial defence mechanisms. This review summarizes some of the most relevant studies in this regard, in order to illuminate this increasingly vast field, while considering the defence mechanisms in relation to the latest advances in phage therapy.

# Innovative strategies in phage therapy: counteracting bacterial defence mechanisms

One of the greatest disadvantages of phage therapy is possibly the frequent and rapid generation of phage-resistant bacterial mutants during treatment. To this regard, the strategy of counteracting these phage defence systems may be a useful way of enhancing the therapeutic outcomes of phages in clinical settings (Figure 2).



**Table 1.** Bacterial defence mechanisms against phages

General mechanism	Specific mechanism	Description	Ref
Adsorption resistance	Modification of bacterial receptor	Either by (i) mutations of cell surface receptors; (ii) inhibitors that compete with the phage; or (iii) alteration of polysaccharide patterns that hide the receptor molecules.	29
	Biofilms	Adsorption of the phage to its bacterial receptor becomes restricted when bacteria form biofilms.	33
	Outer-membrane vesicles (OMVs)	Fragments of the outer membrane of Gram-negative bacteria containing phage receptors may be secreted and act as a bait, leading to phage adsorption and resulting in less successful infections.	34-36
Blocked uptake	Prophages	SIE mechanism: prevention of infection by other phages.	37
·	Small molecules	Anthracyclines, aminoglycosides and viperins: molecules with anti-phage properties, can be secreted by bacteria.	38
Restriction	Restriction modification	Methyltransferase adds methyl groups to the host DNA to distinguish it from the foreign material (especially viral DNA), and the restriction endonuclease recognizes the viral DNA and cuts it.	39
	CRISPR-Cas	Bacteria capture short sequences (protospacers) of invading phages and integrate them in their chromosome (CRISPR array). For subsequent phage infections, the system activates and degrades the phage DNA.	40
Other mechanisms	Abortive infection systems and CBASS	Cells prevent the release of functional virions at the expense of their host cell survival (e.g. inducing programmed cell death).	41,42
	TA	The bacterial toxin induces a global metabolic latency so that phage infection does not progress (no successful phage replication or assembly).	43,44
	QS	Phage adsorption is reduced in the presence of some auto-inducers of the QS network.	45,46

**Table 2.** Phage evasion mechanisms against bacterial defences

Mechanism	Description	Target	References
Alternative adsorption to bacterial surface	Modification of receptor binding protein (RBP) by acquiring mutations that allow recognition of the mutated version of the receptor, or sometimes even a completely different receptor.	Receptor	47
Bacterial capsule degradation	Enzymes like depolymerases allow phages to degrade the extracellular polysaccharides present in the bacterial capsules. In some phages, the RBPs themselves have intrinsic depolymerase activity.	Capsule	48,49
Evasion of Abi	Mutation of specific genes in the phage.	Abi systems	50
Evasion of restriction modification	Reduction in the number or masking of restriction targets incorporating modified bases.	RM systems	51,52
Evasion of TA	Inhibition of a protease that would normally cleave the antitoxin or encoding own antitoxin protein.	TA systems	53,54
Evasion of CRISPR-Cas	(i) Single nucleotide substitution; (ii) complete deletion in the protospacer region or in the conserved motif adjacent to PAM <sup>a</sup> ; (iii) anti-CRISPR systems ( <i>acr-aca</i> ).	CRISPR-Cas systems	55

 $<sup>{}^{\</sup>alpha}\mbox{\rm PAM},$  protospacer adjacent motif.

#### Circumventing mutation of phage receptors

Binding between a phage and its receptor is a critical point in phage infection. However, bacteria have developed defence mechanisms to block phage binding. To circumvent these mechanisms and obtain better results in phage therapy, phage cocktails can be optimized. For example, phage combinations that target different receptors can be prioritized, thereby reducing the likelihood of resistance emergence. For example, Yang et al. designed a cocktail for treating P. aeruginosa that

contained two phages, both with a different receptor, one targeting the O antigen and the other targeting a truncated form of the O antigen that the pathogen produced when it became resistant to the initial O antigen-targeting phage. One of the phage mutants (PaoP5-m1) was found to be excellent for eliminating *P. aeruginosa* mutants with truncated O-antigen structures, as it was able to adsorb and infect all of the strains, regardless of their O-antigen structure. On the other hand, phages may encode enzymes capable of crossing barriers, such as biofilms, capsule layers or the outside of the cell, or the LPSs of the outer

membrane. These proteins are usually located in tail fibres, tail spikes or baseplates. Depolymerases, for example, are a type of enzyme used by phages to access receptors hidden by polysaccharides, or in cases where the polysaccharide chain is the receptor itself, to cleave it, thus stabilizing the binding.<sup>80</sup>

#### Prophage modification

Although phage therapy uses lytic phages, most bacteria have prophages in their genomes, often resulting in a well-documented phage resistance mechanism known as 'superinfection exclusion' (SIE).<sup>37,81</sup> In this situation, a prophage residing in a host cell prevents infection by other similar phage by blocking injection of DNA.<sup>82</sup> However, the presence of prophages in the genome is not always related to SIE defence mechanisms.<sup>81,83,84</sup> Many researchers have claimed that the competitive advantage of lysogens over prophage-free competitors reflects the mutualistic (rather than parasitic) relationship between prophages and bacteria.<sup>85–87</sup>

An interesting approach that paves the way for alternative research aimed at a better understanding of phage therapy is to target these prophages, more specifically using the gene cI, repressor of the lytic cycle, as a therapeutic target. Disruption of the repressor-operator cI by interaction with a small molecule leads to activation of the lytic phase in prophages, thus allowing them to infect their non-lysogenic counterparts and leading to their elimination. <sup>88</sup>

The conversion of a lysogenic phage into a lytic phage displaying activity against multiple clinical isolates of A. baumannii is also an example of the use of engineered phages for therapeutic purposes.<sup>89</sup> In this research, the authors combined the mutant phage with subinhibitory concentrations of different antibiotics and observed a decrease in the frequency of occurrence of phage-resistant bacteria; they also found that combinations including the converted lytic phage increased the survival of infected Galleria mellonella larvae. Alona the same line, a three-phage cocktail composed of engineered phages was administered IV every 12 h for at least 32 weeks to a 15-year-old cystic fibrosis patient with a disseminated Mycobacterium abscessus infection. 74 Importantly, one of these phages (ZoeJ) was lysogenic and converted into a lytic phage by mutation of its repressor gene. This M. abscessus isolate was resistant to all antibiotics tested: clarithromycin, amikacin, tobramycin, ciprofloxacin, moxifloxacin, cefoxitin, co-trimoxazole, doxycycline and linezolid. The phage treatment resulted in a drastic clinical improvement at all levels: sternal wound closure; enhanced liver and lung functions; resolution of infected skin nodules; and weight increase. No adverse effects were observed after this first use of therapeutic phages to treat a human mycobacterial infection.<sup>72</sup>

Prophage induction therapy (a term coined by Lakshminarasimhan in 2022) can potentially be applied to clinical settings to target antibiotic-resistant strains of bacteria. Some researchers have investigated this phenomenon in the gut environment and concluded that medications (including non-antibiotic drugs) inhibiting bacterial growth led to an increase in phage particles due to prophage induction. <sup>90</sup> Importantly, the authors considered this factor an important driver of phage–bacteria dynamics in the gut. One advantage of the prophage induction therapy versus the administration of exogenous lytic phages is that the viral particles

released from the pathogenic bacteria will be highly localized, with lower titres than used for phage therapy, thus minimizing activation of the immune response.<sup>88</sup>

## Clustered regularly interspaced short palindromic repeats (CRISPR-Cas) targeting

CRISPR-Cas systems are the only adaptive immune system observed to date in the prokaryotic world. These are composed of short repeated sequences where spacers become intercalated, conforming the CRISPR-array; this is usually followed by the encoding sequences of CRISPR-associated endonucleases (Cas). Spacer sequences are phage DNA fragments that are cut and integrated into the bacterial CRISPR array, so that bacteria can recognize future infections by phages that they have previously encountered and degrade the DNA of the phages via the Cas endonucleases. 40 The main way that phages evade prokaryotic CRISPR-Cas immunity is by using anti-CRISPR proteins. These basically consist of Acr proteins (typically small proteins of 80–150 amino acids) that inhibit bacterial CRISPR-Cas activity by binding directly to the Cas protein, thereby inactivating it, so that the phages can successfully replicate in the bacterial host. 91 Acr proteins and the Aca (Acr-associated) proteins work via diverse mechanisms to inhibit critical steps of CRISPR immunity, including cas gene expression, 92 assembly of CRISPR ribonucleoprotein complexes, 93,94 recognition of target nucleic acids, 95,96 and recruitment of effector nucleases. 95 Therefore, the use of phages carrying specific acr (or other CRISPR-Cas evasion mechanisms) could be a promising the rapeutic approach against MDR bacteria. and further research is needed in this regard.<sup>9</sup>

#### **Biofilm targeting**

As previously mentioned, biofilm populations are particularly prone to being resistant to antimicrobials and phage attack. Under most conditions, biofilms will allow phage-susceptible bacteria to be protected from phage exposure, if they are growing alongside other cells that are phage resistant; this phenomenon has implications regarding the ecology of phage-bacteria interactions, as well as in the development of phage-based antimicrobial therapeutics. <sup>98</sup> In turn, some phages possess genes coding for extracellular polysaccharide depolymerases that can specifically degrade the polysaccharidic components of biofilms and facilitate the access of phages to deeper layers. <sup>99</sup>

The interaction between phages and bacteria depends on the ability of the phage particles to diffuse through the biofilm, in which cells aggregate and adhere, sometimes hiding phage receptors. <sup>98,100</sup> One interesting example of how a bacteriophage can adapt to biofilm-growing bacteria is the study conducted by Blasco et al. <sup>101</sup> in 2022, in which phages of *A. baumannii* exhibited genomic rearrangement: 10 ORFs were lost and four new ORFs were produced, all of them encoding tail proteins. As a result of this recombination event, a depolymerase-expressing phenotype was visible in 81% of the strains tested. In the same study, a phage cocktail was made with this mutated and adapted phage, together with a phage known to have a depolymerase (B3), and strong activity against 24-h-old biofilms (measured by cfu—enumeration and crystal violet staining) was observed.

Tkhilaishvili *et al.*<sup>102</sup> tested the anti-biofilm activity of a *Staphylococcus aureus*-specific phage, Sb-1, and observed that

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Sb-1 degraded the extracellular matrix formed after 24 h and also targeted persister cells. These researchers assessed the minimum biofilm eradicating concentration (MBEC) and performed confocal laser scanning microscopy (CLSM), which revealed that tail enzymes present in Sb-1 degraded the extracellular polysaccharide component of the matrix of S. aureus ATCC 43300. This is consistent with the findings of Son et al., 103 who reported that a depolymerase present in the lytic S. aureus SAP-2 phage was able to disrupt 48-h-old biofilms [evaluated after safranin staining and scanning electronic microscopy (SEM)]. In another study, Gutiérrez et al. 104 reported that lytic phages were a promising option for fighting biofilm formation in staphylococcal infections. In fact, the results of the study confirmed that lytic phages disrupted 4-h-old biofilms, measured by quantification of cfu and crystal violet staining. With the same pathogen, Alves et al. 105 combined the use of the S. aureus phage K and a newly isolated phage, DRA88, and observed a reduction in growth of a 48-h-old S. aureus biofilm, as observe by crystal violet staining. Rahman et al. 106 combined the phage SAP-26 with erythromycin, vancomycin and rifampicin to treat 24-h-old S. aureus biofilms and obtained a reduction of approximately 28% in the biofilm, as measured by viable cell count, with SAP-26 alone. Erythromycin and vancomycin decreased viable cell counts by 25% and 17%, respectively. However, treatment of the biofilm with the combination of phage and rifampicin vielded a 65% reduction in arowth. The combinations of phage-azithromycin and phage-vancomycin reduced the cell counts by 60% and 40%, respectively, after 24 h.<sup>107</sup>

Another study focusing on the eradication of *in vitro* biofilms using phages is that conducted by Akturk *et al.*, <sup>108</sup> in which they generated dual-species (*S. aureus* and *P. aeruginosa*) 24-h-old biofilms in an *in vitro* artificial dermis model and treated it with specific phages targeting these pathogens (SAFA and EPA1, respectively) and gentamicin. The most effective treatment concerning reduction in viable cells was obtained after multiple doses of EPA1+SAFA+gentamicin, whereas the lowest reduction was produced by application of phages only, followed by another dose of the combined phage treatment.

Many studies have revealed the potential use of phages to eradicate biofilm formation by MDR strains. <sup>107</sup> For example, Khalifa et al. <sup>109</sup> evaluated the lytic activity of an isolated phage from sewage water, EFDG1, against various Enterococcus faecalis and Enterococcus faecium strains, and observed effective activity against planktonic and 2-week-old biofilms, measured by crystal violet staining and CLSM. Importantly, this phage acted on various clinical isolates regardless of their antibiotic resistance profile. In addition, the EFDG1 phage efficiently prevented ex vivo E. faecalis root canal infection, which may be important to prevent persistent infections associated with root canal treatment failure.

Another example is the study conducted by Lehman et al., <sup>110</sup> who evaluated the effect of pretreating hydrogel-coated silicone catheters with mixtures of *P. aeruginosa* and *Proteus mirabilis* phages. The authors used a multiday continuous-flow *in vitro* model with artificial urine medium, in which they produced single- and dual-species biofilms during 96 h. They obtained a 2–4 log reduction in biofilm counts for both species over a period of 48 h. The results of this study suggest that pretreatment of catheters with phage cocktails can significantly reduce mixed-species biofilm formation. In a similar way, Fu et al. <sup>111</sup> showed

the potential of a phage cocktail to prevent *P. aeruginosa* biofilms forming on hydrogel-coated catheters in an *in vitro* model system. The authors observed a reduction in the biofilm viable counts, determined by cfu counts and SEM. Both studies suggest the potential of applying phage cocktails to the surfaces of indwelling medical devices to minimize biofilm formation. <sup>110,111</sup>

Furthermore, Pallavali et al. 112 assessed the effect of lytic phages on 96-h-old biofilms of P. aeruginosa, K. pneumoniae, Escherichia coli and S. aureus, and observed a reduction of nearly 80% in the biomass of biofilms, quantified after cfu enumeration and crystal violet staining. Moreover, Alves et al. 113 showed that a novel bacteriophage cocktail reduced and dispersed P. aeruginosa biofilms under static and flow conditions, therefore having a therapeutic ability to control *P. aeruginosa* infections. For the static model, after contact for 4 h with the phage suspension at an moi of 10, more than 95% of 48-h-old biofilm biomass was eliminated, as measured by crystal violet staining. On the other hand, in the 48-h-old biofilm flow model, slower activity was observed by CLSM; however, 48 h after addition of phage cocktail the biofilm had dispersed. Another example is the case of the phages φMR299-2 and φNH-4, which were able to eliminate the 24-h-old biofilm of P. aeruginosa in vitro (using cystic fibrosis lung airway cells) and in vivo (using a murine lung model). 114,115 Furthermore. Lu et al. 116 reported enhanced dispersion of biofilms with engineered phages, designed from the T7 E. coli phage expressing the dispersin B (DspB). The engineered enzymatic phage reduced bacterial biofilm cell counts by 99.97%.

Other studies have demonstrated the potential of a combined therapy to eradicate biofilms. For instance, the E. coli phage T4 was used in combination with the antibiotic cefotaxime to eradicate 24-h-old F. coli biofilms. The results revealed that the addition of T4 reduced the MBEC of cefotaxime against E. coli biofilms by between 2- and 8-fold, indicating that the combination of T4 and cefotaxime significantly enhanced biofilm eradication.<sup>117</sup> In a similar way, 8-day-old K. pneumoniae biofilms were also eradicated using a combination of phage at an moi of 0.01 with amoxicillin (512 µg/mL). The results of this study showed a significant reduction in the bacterial counts in biofilms after application of combined therapy. 115 In addition, a 12-h-old biofilm of K. pneumoniae was treated by a combination of phage KPO1K2 at an moi of 1 and ciprofloxacin (1 mg/mL), and no significant differences in biofilm removal (quantified by cfu counts) were obtained. However, the combined treatment significantly prevented the emergence of resistance. 118 Other groups used the phage T4 of E. coli and PB-1 of P. aeruginosa in combination with tobramycin at different concentrations (2 and 0.5 µg/mL, respectively) to remove 48-h-old biofilms. The combined treatment led to a 99.99% decrease in the survival of E. coli biofilms measured by cfu/mL count relative to the use of tobramycin alone. However, the combination of tobramycin and P. aeruginosa phage was as effective as tobramycin alone, although the combination reduced the emergence of antibiotic- and phage-resistant cells. <sup>119</sup> Finally, Zhang *et al.* <sup>120</sup> reported that a mixture of RNA phages of P. aeruginosa and chloride reduced the 72-h-old biofilm growth by 94% and removed it in 88% of the cases; the biofilm growth was measured by crystal violet staining.

In general, the main two approaches implemented in regard to the administration of phages to control biofilms are IV bacteriophage therapy and the direct injection of bacteriophages to the site of biofilm on surgical intervention. The first example is documented by Doub *et al.*, <sup>121</sup> but unsuccessful therapeutic outcomes were obtained since the patient continued to have culture-positive *S. aureus* knee infection, suggesting the inability of IV therapy to eradicate the biofilm infection. The second route of administration is documented by Tkhilaishvili *et al.*, <sup>122,123</sup> who treated a chronic relapsing periprosthetic knee infection and chronic osteomyelitis of the femur, caused by an MDR *P. aeruginosa*, with a combined treatment between antibiotics (colistin, meropenem and ceftazidime) and phages, locally applied during surgery.

A systematic review analysing 68 articles on this topic included correlation analysis that revealed some phage parameters relevant to the treatment outcome: higher phage concentrations were strongly associated with better biofilm control; and phages with higher burst sizes and shorter latent periods were the best candidates for controlling biofilms.<sup>124</sup>

The most significant cases of biofilm removal using phages are summarized in Table 4.

#### **Engineered phages**

Phage engineering is also a very important and powerful tool for increasing the likelihood of successful phage therapy; in particular, great effort has been made to extend the host range of phages by genetic engineering, as this is a major limitation of the application of phages in therapy. This strategy could significantly reduce the amount of work needed to search for phages to treat specific bacteria and could lead to the selection of a few well-studied phages that could become 'scaffolds' for the generation of customized phages. Furthermore, this would reduce the variability between treatments and enhance translation to clinical applications. <sup>132</sup>

Yehl et al. 133 targeted regions in the tail fibre of coliphage T3 for mutagenesis, in order to create highly diverse phage libraries that were screened to identify phages with altered host ranges. The authors showed that these engineered phages not only had a broader host range, but they were also able to suppress bacterial resistance to phage infection. They targeted naturally occurring phage-resistant bacterial mutants, which could potentially delay or even prevent the onset of phage resistance. 134 One example of this is a study in which phages containing engineered anti-CRISPR (acr) genes were explored by including type I anti-CRISPR genes (acrIF1, acrIF2 and acrIF3) in the P. aeruginosa phage DMS3/DMS3m to obtain the potential to block bacterial replication and infection. 135 The results indicated that bacterial inhibition required the production of Acrs to be above a specific threshold, so that successful phage replication may be dependent on the competition between CRISPRs and Acrs. This work led to an innovative treatment in which anti-CRISPR phages could be used to treat intractable Pseudomonas disease.

Additionally, natural, engineered and chemically synthesized genomes and re-engineered functional phages that infect Gram-negative bacteria and acid-fast mycobacteria have been assembled and shown to be efficient. 136

A plethora of studies have used genetically modified phages to enhance their potential for biofilm control.<sup>137</sup> Some of these are listed below. For example, expression of enzymes such as

depolymerase or DspB is used to enhance phage activity. <sup>116</sup> Alternatively, insertion of restriction endonuclease genes or modified holin genes or deletion of export protein genes have been explored to minimize inflammatory responses and thus improve phage therapy outcomes. <sup>138–140</sup> One example of the latter is the endosialidases of some coliphages, specialized tail spike proteins that degrade the polysialic acid (polySia) capsule of *E. coli* K1, not only reducing the virulence of the pathogen but, importantly, limiting the tissue damage and the inflammatory processes that occur in response to bacterial invasion. <sup>141</sup>

Simultaneously, Park et al. 142 integrated a CRISPR-Cas system targeting the *nuc* gene (encoding a thermostable nuclease uniquely present in S. aureus, so that the microbiome will not be affected) into the temperate phage \$\phi SaBov\$. They also removed the virulence factors and infected CTH96 S. aureus strains and did not recover any viable cells after treatment with an moi of 100. The authors conducted in vivo studies with a skin infection model in C57BL/6 mice and obtained a reduction in cfu of more than two orders of magnitude after applying \$\phi SaBov-Cas9-nuc embedded into a hydrogel, relative to φSaBov-Cas9-null. To increase the host specificity, the φSaBov tail fibre protein was complemented with that from the phage \$11, which exhibited a broader spectrum, resulting in the specificity being extended to the human pathogenic clones ST1, ST5, ST8 and ST36. 142 Furthermore, this same group studied the effects of \$\phi SaBov-Cas9-nuc on biofilms, both in vitro and in vivo, and enhanced clearance in the biofilm using  $10^8$  pfu/ mL of the CRISPR-Cas-transformed phage φSaBov-Cas9-nuc. 143

Altogether, these findings provide evidence that engineered phages may be a viable alternative approach for use in patients with difficult-to-treat bacterial infections, including those caused by MDR bacteria that do not respond to conventional antibiotic therapy, and also to treat phage-resistant bacteria that have acquired immunity via the CRISPR-Cas system. The most relevant studies concerning engineered phages are summarized in Table 5.

#### **Anti-persister strategies**

Persister cells are a subpopulation of bacteria in a dormant state: they neither grow nor replicate and can re-establish infections once antibiotic stress has been removed.<sup>151</sup>

Persister cells involved in biofilm

The subpopulations of resistant phenotypes within the biofilm have been referred to as persisters. <sup>152</sup>

Phages have two properties that make biofilms susceptible to their action: they produce enzymes that degrade the extracellular matrix; and they can infect persister cells, remaining dormant within them, but reactivating when cells become metabolically active; 125,126 the phages themselves may be the main actors in this reactivation. Furthermore, 'Trojan horse' strategies where phages are used in a first instance to destroy biofilms and activate persister cells, rendering them more susceptible to the antibiotics subsequently introduced, can be considered an elegant pathway to synergy. This strategy takes advantage of the ability of some phages to revert resistance to antibiotics, resensitizing bacteria. 65

Table 3. Clinical case reports focusing on phage therapy

Date and country	Cause of infection	Patient(s)	Pathology/disease	Treatment	Administration route	Clinical outcome	Observation	Reference
2011, Australia P. aeruginosa	ı P. aeruginosa	67-year-old female	Urinary tract infection 6 lytic phage cocktail Pyophage #051007 after was instilled intra-abdominal directly into the resections and bladder every 12 pelvic irradiation for adenocarcinoma	6 lytic phage cocktail	Pyophage #051007 was instilled directly into the bladder every 12 h for 10 days	Resolution of infection No antibiotic- or and microbiological phage-resista improvement arose	No antibiotic- or phage-resistant bacteria arose	8
2016, Bangladesh	E. coli	225 children	Acute bacterial diarrhoea	11 T4-like phage cocktail and Microgen ColiProteus phage	Oral, 4 days	Failure	Overgrowth of Streptococcus gallolyticus and Streptococcus salivarius in the stool	69
2017, USA	MDR A. baumannii 68-year-old	68-year-old	Diabetes, necrotizing pancreatitis (MDR A. baumannii-infected pancreatic pseudocyst)	f 4 anti A. ınii phages	Intracavitary and IV administration	Resolution of the patient's infection	Increased antibiotic sensitivity when phage and antibiotics were simultaneously administered	20
2018, USA	MDR A. <i>baumannii 77-</i> year-old male	77-year-old male	Postoperative craniectomy complication: cerebritis infection	Lytic A. baumannii phage	IV, 98 doses	The patient died	High degree of severity of underlying assault injury	71
2018, Belgium	2018, Belgium XDR <i>P. aeruginosa</i> Male toddler	Male toddler	Sepsis post liver transplantation	Phage-antibiotic combination	IV therapy for 86 days	Complete resolution of all infections, second liver transplantation	Complete resolution of Resistant phage mutants all infections, but not sufficient to second liver re-establish proper transplantation infection	67
2018, USA	Bacteraemia caused by P. aeruginosa	2-year-old boy	DiGeorge syndrome, congenital heart disease	2 lytic phage cocktail IV every 6 h during 36 h	IV every 6 h during 36 h	Death by end-stage cardiac failure	Repeated blood sterilization after each introduction of phage therapy (therapy had to be interrupted)	72
2019, USA	MDR P. aeruginosa 67-year-old and MDR male, Burkholderia 57-year-o dolosa female an 28-year-o female	67-year-old male, 57-year-old female and 28-year-old female	Three lung transplant Cocktails of P. recipients, the third aeruginosa one with cystic phages in fibrosis combinatio antibiotics; phage agai	Cocktails of P. aeruginosa phages in combination with antibiotics; 1 lytic phage against B. dolosa	Nebulized and IV, both antibiotics and phages	Resolution of patient 1 and 2 infections; patient 3 died due to liver and kidney failure	Resistance arose after suspension of phage therapy; challenge in finding lytic phages against <i>B. dolosa</i>	73
2019, USA	M. abscessus	15-year-old	Cystic fibrosis with a disseminated M. abscessus infection	Three-phage cocktail	IV phage treatment, every 12 h for at least 32 weeks	Sternal wound closure, Genome engineering improved liver function, substantial resolution of	Genome engineering	74

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country Cause 2020, USA Mycob	Cause of infection							
		Patient(s)	Pathology/disease	Treatment	Administration route	Clinical outcome	Observation	Reference
						infected skin nodules		
	ınm	56-year-old man Refractory		Muddy phage	IV	Skin lesions improved No resistance observed	No resistance observed	75
che	chelonae		disseminated			and no evidence of		
			cutaneous M.			granulomas		
			chelonae infection					
2022, Belgium PDR K.		30-year-old F	Fracture-related	Phage M1 and	Catheter in place,	Skin graft vascularized, Phage-neutralizing	Phage-neutralizing	92
bue	pneumoniae	bombing	antibiotic-resistant	ceftazidime/	locally	sinus tract closed	antibodies on Days 8-18	
		victim	infection	avibactam,	administered 3	and dry, no signs of	post-phage application	
				tigecycline and	times a day for	infection, gain of	(unlikely to interfere)	
				ciprofloxacin	5 days	muscle		
2023, MDR <i>F</i>	P. aeruginosa 🗓	MDR P. aeruginosa 58-year-old man Infection of		Phages PT07, 14/01, IV 7 days	IV 7 days	Failure	Decrease in the MIC of the	9/
Spain			axillo-bifemoral	PNM and			antimicrobials	
			bypass graft	ceftazidime/				
				avibactam				

PDR, pandrug resistant.

Toxin-antitoxin (TA) targeting

TA modules are generally encoded by two adjacent genes: a stable toxin; and an unstable antitoxin, which is degraded under stress conditions by protease systems, <sup>154</sup> leading to activation of the toxin that often results in reduced bacterial metabolism. <sup>155</sup> One of the main functions of the TA systems is to provide bacteria with another phage resistance mechanism, using the toxin as a general metabolic switch. Therefore, these systems would be a good target for increasing the success of phage therapy. For example, genetically engineered phages that harbour genes encoding toxin inhibitors, or molecules that protect the degradation of the antitoxin by Lon proteases, could potentially be used in phage therapy. <sup>53</sup>

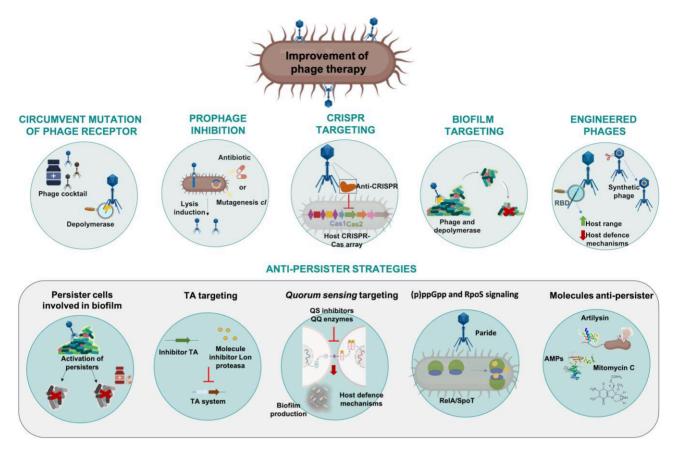
#### Quorum sensing (QS) targeting

QS is bacterial cell-to-cell communication mediated by the production and recognition of small molecules called autoinducers. 156 Several studies clearly show that QS also affects the susceptibility of bacteria to phage infection and the coordination of defence strategies against phages. 45,157-159 Inhibition of QS may be important to enhance phage therapy, as recently demonstrated by Høyland-Kroghsbo et al., 160 who observed overexpression of QS genes in P. aeruginosa infected by the JBD44 phage, demonstrating that QS up-regulates the phage defence systems in the bacterial hosts. The QS network modulates several phage defence mechanisms to appropriately combat infections, shaping the outcomes of phage-host interactions and representing a crucial target to improve phage therapy. In the same line, Shah et al. 161 found that Pseudomonas lysogenic phage DMS3 encoded a QS anti-activator protein. Aas1, which acted as an inhibitor of LasR, the master requlator of QS. The authors also found that Aqs1 protein silences multiple anti-phage defence mechanisms simultaneously, such as Abi systems. Therefore, infection studies with DMS3<sup>ΔAqs1</sup> resulted in 100-fold fewer viable cells, which proved that the presence of Aas1 inhibited Abi-mediated resistance.

Consistently, other studies have linked the inhibition of QS and improved phage infection; for instance, Mion et al. 162 demonstrated the role of the disruption of QS, a strategy known as quorum quenching (QQ), to reduce bacterial virulence and increase both antibiotic and phage treatment efficiency. These researchers used the QQ enzyme SsoPox-W263I, a lactonase able to degrade acyl-homoserine lactones (AHLs), to reduce the virulence and biofilm formation by clinical strains of P. aeruginosa from diabetic foot ulcers, thereby enhancing the susceptibility of phage and antibiotic-sensitive and phage-resistant bacteria to bacteriophages and antibiotics. The study results revealed a reduction of more than 70% in biofilm formation in 6 of 10 strains. Another example is the study by Qin et al., <sup>163</sup> who showed the effect of a QS inhibitor, a penicillic acid, on infection of the strain PAK-AR2 and PAO1 of P. aeruginosa. The results show that supplementation with the QS inhibitor increased the productive infection of the cells by the P. aeruginosa lytic phage C11 and increased the burst size of lytic phage K5 by 45% when cells were infected at the logarithmic phase, relative to the control in the absence of penicillic acid. Finally, the study conducted by Severin et al. 164 demonstrated that QS also up-regulated a different phage defence system in Vibrio cholerae, the cyclic oligonucleotide-based antiphage signalling system (CBASS).

**Fable 3.** Continued





**Figure 2.** Strategies to improve the efficacy of phage therapy, using bacterial defence mechanisms to phage infection as a target. RBD, receptor-binding domain.

Nevertheless, if QS regulates multiple bacterial systems. inhibition of this network could in some cases lead to failure when phages are used as therapeutic agents. For example, Xuan et al. 165 reported that QS up-regulated the expression of phage receptor in P. aeruginosa PAO1 (the O antigen of the LPS), increasing phage adsorption and infection rates. Furthermore, Broniewski et al. 166 showed that a QS inhibitor decreased the phage adsorption rates due to down-regulation of the type IV pilus, which acted as the phage receptor. This caused delayed lysis of bacterial cultures and favoured CRISPR immunity. Therefore, the results of this group suggested that the inhibition of QS may reduce rather than improve the therapeutic efficacy of pilus-specific phages. Similarly, Ghosh et al. 167 demonstrated that a cocktail of different synthetic AHLs or AHL-producing strains led to induction of prophages in E. coli, 168 which could lead to the propagation of these and the subsequent integration in other commensal bacteria. Finally, similar situations have been observed in E. faecalis, V. cholerae and Pseudomonas spp., in which prophage induction was promoted after exposure to autoinducer 2 (AI-2), autoinducer 3,5-dimethylpyrazin-2-ol (DPO) and 2-heptyl-3-hydroxy-4-quinolone (PQS) respectively. 169-174

Overall, prophage induction is a desirable feature in terms of evading phage resistance mediated by prophages (SIE) as well as provoking bacterial lysis; however, caution is required to prevent

horizontal gene transfer and activation of prophage-encoded toxins, among other risks worthy of consideration.<sup>168,175</sup>

These studies confirm the current debate concerning the effects of the QS network on phage defence systems and phage therapy outcomes.

#### (p)ppGpp and RpoS signalling

Most phages enter a more or less stable state of hibernation in deep-dormant host cells (persister cells) and do not produce the lysis of the bacteria cells. However, Maffei et al. 172 found a newly isolated *P. aeruginosa* phage, named Paride, that can directly replicate and induce the lysis of deep-dormant hosts. Efficient replication of Paride on growth-arrested hosts specifically requires cellular stress responses in the form of (p)ppGpp and RpoS signalling that are dispensable for infections of growing hosts. Interestingly, the authors showed that a combination of Paride and meropenem could sterilize deep-dormant cultures *in vitro* and greatly reduced a resilient bacterial infection of a tissue cage implant in mice.

#### Anti-persister molecules

Innovative treatments to tackle dormant, persister bacterial cells, against which antibiotics are not efficient, are needed. In this context, phage therapy could be of benefit. An example

**Table 4.** Treatments and innovation strategies: biofilm targeting

Authors (year)	Mechanisms	Species	References
Pearl et al. (2008) Chegini et al. (2020)	Production of enzymes that degrade extracellular matrix and infect persister cell	P. aeruginosa E. coli	125,126
Eskenazi et al. (2022)	'Trojan horses' strategies where phage was used to destroy biofilm and activate persister cell	K. pneumoniae	65
Dunsing et al. (2019)	Phage carried depolymerase enzyme to counteract the defence of bacteria	Pantoea stewartia	127
Blasco et al. (2022)	Phage cocktail with a host-adapted phage Ab105-2φΔ404ad and phage vB_AbaP_B3 show strong antibiofilm activity	A. baumannii	101
Vidakovic et al. (2018) Chaudhry et al. (2020) Darch et al. (2017)	Phages are retained in biofilms; they confer protection barrier against other bacteria and phages	E. coli P. aeruginosa	128-130
Tkhilaishvili et al. (2017)	Anti-biofilm activity of phage Sb-1; Sb-1 degrades biofilm matrix and target persister cell	S. aureus	102
Son et al. (2010)	Phage SAP-2 carried depolymerase enzyme able to disrupt the biofilm	S. aureus	103
Gutiérrez et al. (2015)	Lytic phages can be efficient biofilm-disrupting agent	Staphylococcus species	104
Khalifa et al. (2015)	Anti-biofilm activity of phage EFDG1	E. faecalis, E. faecium	109
Lehman <i>et al.</i> (2015)	Effect of pretreating hydrogel-coated silicone catheters with phages; these phage cocktails can significantly reduce mixed-species biofilm formation	P. aeruginosa, Proteus mirabilis	110
Alves et al. (2014)	Phage cocktail of phages K and DRA88 reduce biofilm formation	S. aureus	105
Rahman <i>et al.</i> (2011)	Application of phage SAP-26 in combination with different antibiotics against 24-h-old biofilm	S. aureus	106
Akturk et al. (2023)	In vitro dual-species biofilm in combination with gentamicin	P. aeruginosa, S. aureus	131
Fu et al. (2010)	Phage cocktails prevent biofilm formation on catheters in an in vitro model	P. aeruginosa	111
Pallavali et al. (2021)	Lytic phages against 96-h-old multispecies biofilms	P. aeruginosa, K. pneumoniae, E. coli and S. aureus	112
Alves et al. (2016)	Novel phage cocktail reduces and disperses biofilm under static and flow conditions	P. aeruginosa	113
Alemayehu <i>et al.</i> (2012)	Phage φMR299-2 and φNH-4 eliminate <i>P. aeruginosa</i> in murine lung and on cystic fibrosis lung airway cell	P. aeruginosa	114
Lu et al. (2007)	Dispersion of biofilm with 'engineered enzymatic phages'	E. coli	116
Ryan et al. (2012)	Combined therapy to eradicate biofilm: T4 phage with cefotaxime	E. coli	117
Manmeet Sakshi Bedi et al. (2009)	Combined therapy to eradicate biofilm: phages of K. pneumoniae with amoxicillin	K. pneumoniae	115
Verma et al. (2009)	Combined therapy to eradicate 12-h-old biofilms: phage KP01K2 with ciprofloxacin	K. pneumoniae	118
Coulter et al. (2014)	Combined therapy to eradicate 48-h-old biofilms: phage T4 of <i>E. coli</i> and PB-1 of <i>P. aeruginosa</i> with tobramycin	E. coli, P. aeruginosa	119
Zhang et al. (2013)	Mixture of RNA phages and chloride to reduce biofilm	P. aeruginosa	120
Doub et al. (2022)	Recalcitrant MRSA prosthetic knee and femoral lateral plate infection	S. aureus	121
Tkhilaishvili et al. (2019)	Direct phage injection in a periprosthetic joint infection and osteomyelitis	P. aeruginosa	122

of the use of lytic phages against an imipenem-persister clinical isolate of *K. pneumoniae* has been reported by Pacios et al. <sup>173</sup> in a study in which a lytic phage was used in combination with the repurposed anticancer drug mitomycin C (also considered a natural antibiotic) and the conventional imipenem. Both combinations resulted in the death of persister cells and decreased the emergence of *in vitro* resistant mutants. These results were confirmed in the *in vivo G. mellonella* model, in which the combination significantly reduced the mortality rate of the larvae. Briers et al. <sup>174</sup> also described a novel artilysin (an outer membrane-penetrating, phage-derived)

endolysin), named Art-175, that is able to rapidly and effectively pass through the outer membrane of *P. aeruginosa* persister cells and exert its bactericidal activity within the cell, without the need for any metabolic activity. Endolysins are peptidoglycan-degrading enzymes synthesized by phages, used at the end of their replication cycle to hydrolyse the peptidoglycan from within and allow the release of newly formed virions. <sup>33–35,175</sup>

All of these studies provide some insights into the treatment of persister cells, prone to forming biofilms that are difficult to eradicate.



**Table 5.** Report cases of engineered phages

Authors (year)	Target	Methods	References
Pei and Lamas-Samanamud (2014)	T7 phage was constructed to encode a lactonase enzyme for quenching of QS	Phage-displayed vector	144
Park et al. (2017)	Temperate phage φSaBov of S. aureus	CRISPR-Cas9 genomic editing technique	142
Yehl <i>et al.</i> (2019)	A region in the tail fibre of coliphage T3	Mutagenesis	133
Blasco et al. (2019)	Repressor gene cI of lysogenic phage of A. baumannii	Knockout by homologous recombination	89
Dedrick et al. (2019)	Repressor gene of lysogenic phages of M. abscessus	Phage recombineering of electroporated DNA (BRED)	74
Pires <i>et al.</i> (2021)	Reduction of genome encoding hypothetical proteins (48%)	Yeast-based phage-engineering platform	145
Qin et al. (2022)	Introducing type I anti-CRISPR genes (AcrIF1, AcrIF2 and AcrIF3) into the <i>P. aeruginosa</i> phage DMS3/DMS3m	Homologous recombination	135
Mitsunaka <i>et al.</i> (2022)	Deleting the c2 lysis-repressor gene in P22 Salmonella phage, host-range increase of T3 coliphage with T7 tail fibre, engineering of mycophage D29 to show luminescence (addition of the Nluc gene)	In vitro genome assembly and installation into appropriate host bacteria or a cell-free transcription and translation (TXTL) system	136
Nick et al. (2022)	Repressor gene 45 of lytic phage of <i>M. abscessus</i>	Phage recombineering of electroporated DNA (BRED)	146
Emslander et al. (2022)	E. coli, Yersinia pestis and K. pneumoniae	Cell-free method and proteomic characterization (MS)	147
Liyanagedera <i>et al.</i> (2022)	Incorporate a SpyTag moiety on the capsid head to enable rapid post-synthetic modification	Scaffold phage	148
Fa-arun <i>et al.</i> (2023)	Phage tail for the delivery of a Cas9 antimicrobial into clinically relevant human gut pathogens <i>Shigella flexneri</i> and <i>E. coli</i> 0157:H7	Cosmid system	149
Prokopczuk et al. (2023)	Repressor and the excisionase genes of Pf4 phage in <i>P. aeruginosa</i> PAO1 strain	Mutagenesis	150

#### **Conclusions**

Because of the constant adaptation over hundreds of millions of years of co-evolution, bacteria and phages have acquired a plethora of mechanisms to defend themselves against infection and to neutralize these defence systems. This review summarizes some of the most relevant bacterial and phage defence mechanisms in order to throw some light on this increasingly vast topic, and it outlines the latest advances in phage therapy with particular focus on targeting the main bacterial defence systems (mutation of phage receptors, prophages, CRISPR-Cas, biofilm, TA systems, QS). The latter two are involved in the development of persister cells. It is therefore essential to develop more antipersister strategies to counteract these bacterial phenotypes. In this context, phage engineering is of particular interest to improve the potential use of personalized phages in therapy.

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#### **Transparency declarations**

None to declare.

#### References

- **1** Antimicrobial Resistance Collaborators. Global burden of bacterial antimicrobial resistance in 2019: a systematic analysis. *Lancet* 2022; **399**: 629–55. https://doi.org/10.1016/S0140-6736(21)02724-0
- **2** Wright GD, Poinar H. Antibiotic resistance is ancient: implications for drug discovery. *Trends Microbiol* 2012; **20**: 157–9. https://doi.org/10.1016/j.tim.2012.01.002

- D'Costa VM, King CE, Kalan L *et al.* Antibiotic resistance is ancient. *Nature* 2011; **477**: 457–61. https://doi.org/10.1038/nature10388
- Bartlett JG, Gilbert DN, Spellberg B. Seven ways to preserve the miracle of antibiotics. *Clin Infect Dis* 2013; **56**: 1445–50. https://doi.org/10.1093/cid/cit070
- Trastoy R, Manso T, Fernandez-Garcia L *et al.* Mechanisms of bacterial tolerance and persistence in the gastrointestinal and respiratory environments. *Clin Microbiol Rev* 2018; **31**: e00023-18. https://doi.org/10.1128/CMR.00023-18
- Gu D, Dong N, Zheng Z *et al.* A fatal outbreak of ST11 carbapenem-resistant hypervirulent *Klebsiella pneumoniae* in a Chinese hospital: a molecular epidemiological study. *Lancet Infect Dis* 2018; **18**: 37–46. https://doi.org/10.1016/S1473-3099(17)30489-9
- Zeng L, Yang C, Zhang J *et al.* An outbreak of carbapenem-resistant *Klebsiella pneumoniae* in an intensive care unit of a major teaching hospital in Chongqing, China. *Front Cell Infect Microbiol* 2021; **11**: 656070. https://doi.org/10.3389/fcimb.2021.656070
- Du H, Chen L, Tang YW *et al.* Emergence of the *mcr-1* colistin resistance gene in carbapenem-resistant Enterobacteriaceae. *Lancet Infect Dis* 2016; **16**: 287–8. https://doi.org/10.1016/S1473-3099(16)00056-6
- **9** Trongjit S, Assavacheep P, Samngamnim S *et al.* Plasmid-mediated colistin resistance and ESBL production in *Escherichia coli* from clinically healthy and sick pigs. *Sci Rep* 2022; **12**: 2466. https://doi.org/10.1038/s41598-022-06415-0
- Majewski P, Gutowska A, Smith DGE *et al.* Plasmid mediated *mcr-1.1* colistin-resistance in clinical extraintestinal *Escherichia coli* strains isolated in Poland. *Front Microbiol* 2021; **12**: 547020. https://doi.org/10.3389/fmicb.2021.547020
- Witney AA, Gould KA, Pope CF *et al.* Genome sequencing and characterization of an extensively drug-resistant sequence type 111 serotype 012 hospital outbreak strain of *Pseudomonas aeruginosa. Clin Microbiol Infect* 2014; **20**: O609–18. https://doi.org/10.1111/1469-0691.12528
- Thoma R, Seneghini M, Seiffert SN *et al.* The challenge of preventing and containing outbreaks of multidrug-resistant organisms and *Candida auris* during the coronavirus disease 2019 pandemic: report of a carbapenem-resistant *Acinetobacter baumannii* outbreak and a systematic review of the literature. *Antimicrob Resist Infect Control* 2022; **11**: 12. https://doi.org/10.1186/s13756-022-01052-8
- Diniz Rocha VF, Cavalcanti TP, Azevedo J et al. Outbreak of Stenotrophomonas maltophilia and Burkholderia cepacia bloodstream infections at a hemodialysis center. Am J Trop Med Hyg 2020; **104**: 848–53. https://doi.org/10.4269/ajtmh.20-1035
- Guyot A, Turton JF, Garner D. Outbreak of *Stenotrophomonas maltophilia* on an intensive care unit. *J Hosp Infect* 2013; **85**: 303–7. https://doi.org/10.1016/j.jhin.2013.09.007
- Menekşe Ş, Tanrıverdi ES, Oğuş H *et al. Stenotrophomonas maltophilia* outbreak with a commercial blood gas injector as the culprit and interventions for source and prevention: a possible passage between patient and ECMO water heater device. *Am J Infect Control* 2023; **51**: 533–8. https://doi.org/10.1016/j.ajic.2022.07.012
- Möllers M, von Wahlde MK, Schuler F *et al.* Outbreak of MRSA in a gynecology/obstetrics department during the COVID-19 pandemic: a cautionary tale. *Microorganisms* 2022; **10**: 689. https://doi.org/10.3390/microorganisms10040689
- Rubin IM, Hansen TA, Klingenberg AM *et al.* A sporadic four-year hospital outbreak of a ST97-IVa MRSA with half of the patients first identified in the community. *Front Microbiol* 2018; **9**: 1494. https://doi.org/10.3389/fmicb.2018.01494
- Hutchings MI, Truman AW, Wilkinson B. Antibiotics: past, present and future. *Curr Opin Microbiol* 2019; **51**: 72–80. https://doi.org/10.1016/j.mib. 2019.10.008

- Ventola CL. The antibiotic resistance crisis: part 1: causes and threats. *PT* 2015; **40**: 277–83.
- Qi YK, Tang X, Wei NN *et al.* Discovery, synthesis, and optimization of teixobactin, a novel antibiotic without detectable bacterial resistance. *J Pept Sci* 2022; **28**: e3428. https://doi.org/10.1002/psc.3428
- Gunjal VB, Thakare R, Chopra S *et al.* Teixobactin: a paving stone toward a new class of antibiotics? *J Med Chem* 2020; **63**: 12171–95. https://doi.org/10.1021/acs.jmedchem.0c00173
- **22** Lloyd DG, Schofield BJ, Goddard MR *et al. De novo* resistance to Arg<sub>10</sub>-teixobactin occurs slowly and is costly. *Antimicrob Agents Chemother* 2020; **65**: e01152-20. https://doi.org/10.1128/AAC.01152-20
- Pacios O, Blasco L, Bleriot I *et al.* Strategies to combat multidrug-resistant and persistent infectious diseases. *Antibiotics* (*Basel*) 2020; **9**: 65. https://doi.org/10.3390/antibiotics9020065
- **24** Parmanik A, Das S, Kar B *et al.* Current treatment strategies against multidrug-resistant bacteria: a review. *Curr Microbiol* 2022; **79**: 388. https://doi.org/10.1007/s00284-022-03061-7
- Bergh O, Børsheim KY, Bratbak G *et al.* High abundance of viruses found in aquatic environments. *Nature* 1989; **340**: 467–8. https://doi.org/10.1038/340467a0
- Mäntynen S, Laanto E, Oksanen HM *et al.* Black box of phage-bacterium interactions: exploring alternative phage infection strategies. *Open Biol* 2021; **11**: 210188. https://doi.org/10.1098/rsob.210188
- Howard-Varona C, Hargreaves KR, Abedon ST *et al.* Lysogeny in nature: mechanisms, impact and ecology of temperate phages. *ISME J* 2017; **11**: 1511–20. https://doi.org/10.1038/ismej.2017.16
- Du Toit A. Viral infection: the language of phages. *Nat Rev Microbiol* 2017; **15**: 134–5. https://doi.org/10.1038/nrmicro.2017.8
- Labrie SJ, Samson JE, Moineau S. Bacteriophage resistance mechanisms. *Nat Rev Microbiol* 2010; **8**: 317–27. https://doi.org/10.1038/nrmicro2315
- Samson JE, Magadán AH, Sabri M *et al.* Revenge of the phages: defeating bacterial defences. *Nat Rev Microbiol* 2013; **11**: 675–87. https://doi.org/10.1038/nrmicro3096
- Ambroa A, Blasco L, López M *et al*. Genomic analysis of molecular bacterial mechanisms of resistance to phage infection. *Front Microbiol* 2022; **12**: 784949. https://doi.org/10.3389/fmicb.2021.784949
- Bleriot I, Blasco L, Pacios O *et al.* Proteomic study of the interactions between phages and the bacterial host *Klebsiella pneumoniae*. *Microbiol Spectr* 2023; **11**: e0397422. https://doi.org/10.1128/spectrum. 03974-22
- Visnapuu A, Van der Gucht M, Wagemans J et al. Deconstructing the phage-bacterial biofilm interaction as a basis to establish new antibiofilm strategies. *Viruses* 2022; **14**: 1057. https://doi.org/10.3390/v14051057
- Reyes-Robles T, Dillard RS, Cairns LS *et al. Vibrio cholerae* outer membrane vesicles inhibit bacteriophage infection. *J Bacteriol* 2018; **200**: e00792-17. https://doi.org/10.1128/JB.00792-17
- Stephan MS, Broeker NK, Saragliadis A *et al.* Analysis of O-antigen-specific bacteriophage P22 inactivation by *Salmonella* outer membrane vesicles. *Front Microbiol* 2020; **11**: 510638. https://doi.org/10.3389/fmicb.2020.510638
- Yasuda M, Yamamoto T, Nagakubo T *et al.* Phage genes induce quorum sensing signal release through membrane vesicle formation. *Microbes Environ* 2022; **37**: ME21067. https://doi.org/10.1264/jsme2.ME21067
- Susskind MM, Wright A, Botstein D. Superinfection exclusion by P22 prophage in lysogens of *Salmonella typhimurium*. IV. Genetics and physiology of *sieB* exclusion. *Virology* 1974; **62**: 367–84. https://doi.org/10.1016/0042-6822(74)90399-7
- **38** Hardy A, Kever L, Frunzke J. Antiphage small molecules produced by bacteria—beyond protein-mediated defenses. *Trends Microbiol* 2023; **31**: 92–106. https://doi.org/10.1016/j.tim.2022.08.001

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- Rusinov IS, Ershova AS, Karyagina AS *et al.* Avoidance of recognition sites of restriction-modification systems is a widespread but not universal anti-restriction strategy of prokaryotic viruses. *BMC Genomics* 2018; **19**: 885. https://doi.org/10.1186/s12864-018-5324-3
- Barrangou R, Fremaux C, Deveau H *et al.* CRISPR provides acquired resistance against viruses in prokaryotes. *Science* 2007; **315**: 1709–12. https://doi.org/10.1126/science.1138140
- Aframian N, Eldar A. Abortive infection antiphage defense systems: separating mechanism and phenotype. *Trends Microbiol* 2023; **31**: 1003–12. https://doi.org/10.1016/j.tim.2023.05.002
- Lopatina A, Tal N, Sorek R. Abortive infection: bacterial suicide as an antiviral immune strategy. *Annu Rev Virol* 2020; **7**: 371–84. https://doi.org/10.1146/annurev-virology-011620-040628
- Bleriot I, Blasco L, Pacios O *et al*. The role of PemIK (PemK/PemI) type II TA system from *Klebsiella pneumoniae* clinical strains in lytic phage infection. *Sci Rep* 2022; **12**: 4488. https://doi.org/10.1038/s41598-022-08111-5
- Hazan R, Engelberg-Kulka H. *Escherichia coli mazEF*-mediated cell death as a defense mechanism that inhibits the spread of phage P1. *Mol Genet Genomics* 2004; **272**: 227–34. https://doi.org/10.1007/s00438-004-1048-y
- Tan D, Svenningsen SL, Middelboe M. Quorum sensing determines the choice of antiphage defense strategy in *Vibrio anguillarum. mBio* 2015; **6**: e00627. https://doi.org/10.1128/mBio.00627-15
- Hoque MM, Naser IB, Bari SM *et al.* Quorum regulated resistance of *Vibrio cholerae* against environmental bacteriophages. *Sci Rep* 2016; **6**: 37956. https://doi.org/10.1038/srep37956
- Meyer JR, Dobias DT, Weitz JS *et al.* Repeatability and contingency in the evolution of a key innovation in phage lambda. *Science* 2012; **335**: 428–32. https://doi.org/10.1126/science.1214449
- **48** Scholl D, Adhya S, Merril C. *Escherichia coli* K1's capsule is a barrier to bacteriophage T7. *Appl Environ Microbiol* 2005; **71**: 4872–4. https://doi.org/10.1128/AEM.71.8.4872-4874.2005
- **49** Cornelissen A, Ceyssens PJ, Krylov VN et al. Identification of EPS-degrading activity within the tail spikes of the novel *Pseudomonas putida* phage AF. *Virology* 2012; **434**: 251–6. https://doi.org/10.1016/j. virol.2012.09.030
- Labrie SJ, Tremblay DM, Moisan M *et al.* Involvement of the major capsid protein and two early-expressed phage genes in the activity of the lactococcal abortive infection mechanism AbiT. *Appl Environ Microbiol* 2012; **78**: 6890-9. https://doi.org/10.1128/AEM.01755-12
- **51** Iida S, Streiff MB, Bickle TA *et al.* Two DNA antirestriction systems of bacteriophage P1, *darA*, and *darB*: characterization of *darA*<sup>-</sup> phages. *Virology* 1987; **157**: 156–66. https://doi.org/10.1016/0042-6822(87)90324-2
- Murphy J, Mahony J, Ainsworth S *et al.* Bacteriophage orphan DNA methyltransferases: insights from their bacterial origin, function, and occurrence. *Appl Environ Microbiol* 2013; **79**: 7547–55. https://doi.org/10.1128/AEM.02229-13
- Sberro H, Leavitt A, Kiro R *et al.* Discovery of functional toxin/antitoxin systems in bacteria by shotgun cloning. *Mol Cell* 2013; **50**: 136–48. https://doi.org/10.1016/j.molcel.2013.02.002
- Otsuka Y, Yonesaki T. Dmd of bacteriophage T4 functions as an antitoxin against *Escherichia coli* LsoA and RnlA toxins. *Mol Microbiol* 2012; **83**: 669–81. https://doi.org/10.1111/j.1365-2958.2012.07975.x
- Deveau H, Barrangou R, Garneau JE *et al.* Phage response to CRISPR-encoded resistance in *Streptococcus thermophilus*. *J Bacteriol* 2008; **190**: 1390–400. https://doi.org/10.1128/JB.01412-07
- Young R, Gill JJ. Phage therapy redux—what is to be done? *Science* 2015; **350**: 1163–4. https://doi.org/10.1126/science.aad6791
- Ferry T, Kolenda C, Briot T *et al*. Past and future of phage therapy and phage-derived proteins in patients with bone and joint infection. *Viruses* 2021; **13**: 2414. https://doi.org/10.3390/v13122414

- Uyttebroek S, Chen B, Onsea J *et al.* Safety and efficacy of phage therapy in difficult-to-treat infections: a systematic review. *Lancet Infect Dis* 2022; **22**: e208–20. https://doi.org/10.1016/S1473-3099(21)00612-5
- Tamma PD, Souli M, Billard M et al. Safety and microbiological activity of phage therapy in persons with cystic fibrosis colonized with *Pseudomonas aeruginosa*: study protocol for a phase 1b/2, multicenter, randomized, double-blind, placebo-controlled trial. *Trials* 2022; **23**: 1057. https://doi.org/10.1186/s13063-022-07047-5
- Gómez-Ochoa SA, Pitton M, Valente LG *et al*. Efficacy of phage therapy in preclinical models of bacterial infection: a systematic review and meta-analysis. *Lancet Microbe* 2022; **3**: e956–68. https://doi.org/10.1016/S2666-5247(22)00288-9
- Steele A, Stacey HJ, de Soir S *et al.* The safety and efficacy of phage therapy for superficial bacterial infections: a systematic review. *Antibiotics* (*Basel*) 2020; **9**: 754. https://doi.org/10.3390/antibiotics9110754
- **62** Pirnay JP, Djebara S, Steurs G *et al.* Retrospective, observational analysis of the first one hundred consecutive cases of personalized bacteriophage therapy of difficult-to-treat infections facilitated by a Belgian consortium. *medRxiv* 2023; https://doi.org/10.1101/2023.08.28.23294728
- Green SI, Clark JR, Santos HH *et al.* A retrospective, observational study of 12 cases of expanded access customized phage therapy: production, characteristics, and clinical outcomes. *Clin Infect Dis* 2023; **77**: 1079–91. https://doi.org/10.1093/cid/ciad335
- **64** Comeau AM, Tétart F, Trojet SN *et al.* Phage-antibiotic synergy (PAS):  $\beta$ -lactam and quinolone antibiotics stimulate virulent phage growth. *PLoS One* 2007; **2**: e799. https://doi.org/10.1371/journal.pone.0000799
- **65** Eskenazi A, Lood C, Wubbolts J *et al.* Combination of pre-adapted bacteriophage therapy and antibiotics for treatment of fracture-related infection due to pandrug-resistant *Klebsiella pneumoniae. Nat Commun* 2022; **13**: 302. https://doi.org/10.1038/s41467-021-27656-z
- Burrowes BH, Molineux IJ, Fralick JA. Directed in vitro evolution of therapeutic bacteriophages: the Appelmans protocol. *Viruses* 2019; **11**: 241. https://doi.org/10.3390/v11030241
- **67** Van Nieuwenhuyse B, Van der Linden D, Chatzis O *et al.* Bacteriophage-antibiotic combination therapy against extensively drug-resistant *Pseudomonas aeruginosa* infection to allow liver transplantation in a toddler. *Nat Commun* 2022; **13**: 5725. https://doi.org/10.1038/s41467-022-33294-w
- Khawaldeh A, Morales S, Dillon B *et al.* Bacteriophage therapy for refractory *Pseudomonas aeruginosa* urinary tract infection. *J Med Microbiol* 2011; **60**: 1697–700. https://doi.org/10.1099/jmm.0.029744-0
- Sarker SA, Sultana S, Reuteler G *et al.* Oral phage therapy of acute bacterial diarrhea with two coliphage preparations: a randomized trial in children from Bangladesh. *EBioMedicine* 2016; **4**: 124–37. https://doi.org/10.1016/j.ebiom.2015.12.023
- **70** Schooley RT, Biswas B, Gill JJ *et al.* Development and use of personalized bacteriophage-based therapeutic cocktails to treat a patient with a disseminated resistant *Acinetobacter baumannii* infection. *Antimicrob Agents Chemother* 2017; **61**: e00954-17. https://doi.org/10.1128/AAC. 00954-17
- LaVergne S, Hamilton T, Biswas B *et al.* Phage therapy for a multidrug-resistant *Acinetobacter baumannii* craniectomy site infection. *Open Forum Infect Dis* 2018; **5**: ofy064. https://doi.org/10.1093/ofid/ofy064
- Duplessis C, Biswas B, Hanisch B *et al.* Refractory *Pseudomonas* bacteremia in a 2-year-old sterilized by bacteriophage therapy. *J Pediatric Infect Dis Soc* 2018; **7**: 253–6. https://doi.org/10.1093/jpids/pix056
- Aslam S, Courtwright AM, Koval C *et al*. Early clinical experience of bacteriophage therapy in 3 lung transplant recipients. *Am J Transplant* 2019; **19**: 2631–9. https://doi.org/10.1111/ajt.15503
- Dedrick RM, Guerrero-Bustamante CA, Garlena RA *et al.* Engineered bacteriophages for treatment of a patient with a disseminated

- drug-resistant *Mycobacterium abscessus*. *Nat Med* 2019; **25**: 730–3. https://doi.org/10.1038/s41591-019-0437-z
- Little JS, Dedrick RM, Freeman KG *et al.* Bacteriophage treatment of disseminated cutaneous *Mycobacterium chelonae* infection. *Nat Commun* 2022; **13**: 2313. https://doi.org/10.1038/s41467-022-29689-4
- **76** Blasco L, López-Hernández I, Rodríguez-Fernández M *et al.* Case report: analysis of phage therapy failure in a patient with a *Pseudomonas aeruginosa* prosthetic vascular graft infection. *Front Med (Lausanne)* 2023; **10**: 1199657. https://doi.org/10.3389/fmed.2023.1199657
- Loc-Carrillo C, Abedon ST. Pros and cons of phage therapy. *Bacteriophage* 2011; **1**: 111–4. https://doi.org/10.4161/bact.1.2.14590
- Gordillo Altamirano FL, Barr JJ. Unlocking the next generation of phage therapy: the key is in the receptors. *Curr Opin Biotechnol* 2021; **68**: 115–23. https://doi.org/10.1016/j.copbio.2020.10.002
- Yang Y, Shen W, Zhong Q *et al.* Development of a bacteriophage cocktail to constrain the emergence of phage-resistant *Pseudomonas aeruginosa. Front Microbiol* 2020; **11**: 327. https://doi.org/10.3389/fmicb.2020.00327
- Knecht LE, Veljkovic M, Fieseler L. Diversity and function of phage encoded depolymerases. *Front Microbiol* 2020; **10**: 2949. https://doi.org/10. 3389/fmich 2019 02949
- Bleriot I, Trastoy R, Blasco L *et al.* Genomic analysis of 40 prophages located in the genomes of 16 carbapenemase-producing clinical strains of *Klebsiella pneumoniae. Microb Genom* 2020; **6**: e000369. https://doi.org/10.1099/mgen.0.000369
- Donnelly-Wu MK, Jacobs WR, Hatfull GF. Superinfection immunity of mycobacteriophage L5: applications for genetic transformation of mycobacteria. *Mol Microbiol* 1993; **7**: 407–17. https://doi.org/10.1111/j.1365-2958.1993.tb01132.x
- Hao G, Shu R, Ding L *et al.* Bacteriophage SRD2021 recognizing capsular polysaccharide shows therapeutic potential in serotype K47 *Klebsiella pneumoniae* infections. *Antibiotics (Basel)* 2021; **10**: 894. https://doi.org/10.3390/antibiotics10080894
- Serian D, Churin Y, Hammerl JA et al. Characterization of temperate LPS-binding *Bordetella avium* phages that lack superinfection immunity. *Microbiol Spectr* 2023; **11**: e0370222. https://doi.org/10.1128/spectrum. 03702-22
- Bondy-Denomy J, Davidson AR. When a virus is not a parasite: the beneficial effects of prophages on bacterial fitness. *J Microbiol* 2014; **52**: 235–42. https://doi.org/10.1007/s12275-014-4083-3
- **86** Nanda AM, Thormann K, Frunzke J. Impact of spontaneous prophage induction on the fitness of bacterial populations and host-microbe interactions. *J Bacteriol* 2015; **197**: 410–9. https://doi.org/10.1128/JB.02230-14
- Fillol-Salom A, Alsaadi A, Sousa JAM *et al.* Bacteriophages benefit from generalized transduction. *PLoS Pathog* 2019; **15**: e1007888. https://doi.org/10.1371/journal.ppat.1007888
- Lakshminarasimhan A. Prophage induction therapy: activation of the lytic phase in prophages for the elimination of pathogenic bacteria. *Med Hypothesis* 2022; **169**: 110980. https://doi.org/10.1016/j.mehy.2022.110980
- Blasco L, Ambroa A, Lopez M *et al.* Combined use of the Ab105-2phiDeltaCI lytic mutant phage and different antibiotics in clinical isolates of multi-resistant *Acinetobacter baumannii. Microorganisms* 2019; **7**: 556. https://doi.org/10.3390/microorganisms7110556
- Sutcliffe SG, Shamash M, Hynes AP *et al.* Common oral medications lead to prophage induction in bacterial isolates from the human gut. *Viruses* 2021; **13**: 455. https://doi.org/10.3390/v13030455
- Li Y, Bondy-Denomy J. Anti-CRISPRs go viral: the infection biology of CRISPR-Cas inhibitors. *Cell Host Microbe* 2021; **29**: 704–14. https://doi.org/10.1016/j.chom.2020.12.007
- Osuna BA, Karambelkar S, Mahendra C *et al. Listeria* phages induce Cas9 degradation to protect lysogenic genomes. *Cell Host Microbe* 2020; **28**: 31–40.e9. https://doi.org/10.1016/j.chom.2020.04.001

- **93** Mahendra C, Christie KA, Osuna BA *et al.* Broad-spectrum anti-CRISPR proteins facilitate horizontal gene transfer. *Nat Microbiol* 2020; **5**: 620–9. https://doi.org/10.1038/s41564-020-0692-2
- Knott GJ, Thornton BW, Lobba MJ *et al.* Broad-spectrum enzymatic inhibition of CRISPR-Cas12a. *Nat Struct Mol Biol* 2019; **26**: 315–21. https://doi.org/10.1038/s41594-019-0208-z
- Bondy-Denomy J, Garcia B, Strum S *et al.* Multiple mechanisms for CRISPR-Cas inhibition by anti-CRISPR proteins. *Nature* 2015; **526**: 136–9. https://doi.org/10.1038/nature15254
- Meeske AJ, Jia N, Cassel AK *et al.* A phage-encoded anti-CRISPR enables complete evasion of type VI-A CRISPR-Cas immunity. *Science* 2020; **369**: 54–9. https://doi.org/10.1126/science.abb6151
- Vyas P, Harish. Anti-CRISPR proteins as a therapeutic agent against drug-resistant bacteria. *Microbiol Res* 2022; **257**: 126963. https://doi.org/10.1016/j.micres.2022.126963
- Simmons EL, Bond MC, Koskella B *et al.* Biofilm structure promotes coexistence of phage-resistant and phage-susceptible bacteria. *mSystems* 2020; **5**: e00877-19. https://doi.org/10.1128/mSystems.00877-19
- Liu S, Lu H, Zhang S *et al.* Phages against pathogenic bacterial biofilms and biofilm-based infections: a review. *Pharmaceutics* 2022; **14**: 427. https://doi.org/10.3390/pharmaceutics14020427
- Heilmann S, Sneppen K, Krishna S. Coexistence of phage and bacteria on the boundary of self-organized refuges. *Proc Natl Acad Sci U S A* 2012; **109**: 12828–33. https://doi.org/10.1073/pnas.1200771109
- Blasco L, Bleriot I, de Aledo MG *et al.* Development of an anti-*Acinetobacter baumannii* biofilm phage cocktail: genomic adaptation to the host. *Antimicrob Agents Chemother* 2022; **66**: e0192321. https://doi.org/10.1128/aac.01923-21
- Tkhilaishvili T, Lombardi L, Klatt AB *et al.* Bacteriophage Sb-1 enhances antibiotic activity against biofilm, degrades exopolysaccharide matrix and targets persisters of *Staphylococcus aureus*. *Int J Antimicrob Agents* 2018; **52**: 842–53. https://doi.org/10.1016/j.ijantimicag.2018.09.006
- Son JS, Lee SJ, Jun SY *et al*. Antibacterial and biofilm removal activity of a podoviridae *Staphylococcus aureus* bacteriophage SAP-2 and a derived recombinant cell-wall-degrading enzyme. *Appl Microbiol Biotechnol* 2010; **86**: 1439–49. https://doi.org/10.1007/s00253-009-2386-9
- Gutiérrez D, Vandenheuvel D, Martínez B *et al.* Two phages, phiIPLA-RODI and phiIPLA-C1C, lyse mono- and dual-species staphylococcal biofilms. *Appl Environ Microbiol* 2015; **81**: 3336–48. https://doi.org/10.1128/AEM.03560-14
- **105** Alves DR, Gaudion A, Bean JE *et al.* Combined use of bacteriophage K and a novel bacteriophage to reduce *Staphylococcus aureus* biofilm formation. *Appl Environ Microbiol* 2014; **80**: 6694–703. https://doi.org/10.1128/AEM.01789-14
- Rahman M, Kim S, Kim SM *et al.* Characterization of induced *Staphylococcus aureus* bacteriophage SAP-26 and its anti-biofilm activity with rifampicin. *Biofouling* 2011; **27**: 1087-93. https://doi.org/10.1080/08927014.2011.631169
- Pires DP, Melo L, Vilas Boas D *et al.* Phage therapy as an alternative or complementary strategy to prevent and control biofilm-related infections. *Curr Opin Microbiol* 2017; **39**: 48–56. https://doi.org/10.1016/j.mib. 2017.09.004
- Akturk E, Melo LDR, Oliveira H *et al.* Combining phages and antibiotic to enhance antibiofilm efficacy against an *in vitro* dual species wound biofilm. *Biofilm* 2023; **6**: 100147. https://doi.org/10.1016/j.bioflm.2023.100147
- Khalifa L, Brosh Y, Gelman D *et al.* Targeting *Enterococcus faecalis* biofilms with phage therapy. *Appl Environ Microbiol* 2015; **81**: 2696–705. https://doi.org/10.1128/AEM.00096-15
- **110** Lehman SM, Donlan RM. Bacteriophage-mediated control of a two-species biofilm formed by microorganisms causing catheter-associated urinary tract infections in an *in vitro* urinary catheter model.

JAR

Antimicrob Agents Chemother 2015; **59**: 1127–37. https://doi.org/10. 1128/AAC.03786-14

- **111** Fu W, Forster T, Mayer O *et al.* Bacteriophage cocktail for the prevention of biofilm formation by *Pseudomonas aeruginosa* on catheters in an *in vitro* model system. *Antimicrob Agents Chemother* 2010; **54**: 397–404. https://doi.org/10.1128/AAC.00669-09
- Pallavali RR, Degati VL, Narala VR *et al.* Lytic bacteriophages against bacterial biofilms formed by multidrug-resistant *Pseudomonas aeruginosa, Escherichia coli, Klebsiella pneumoniae,* and *Staphylococcus aureus* isolated from burn wounds. *Phage (New Rochelle)* 2021; **2**: 120–30. https://doi.org/10.1089/phage.2021.0004
- Alves DR, Perez-Esteban P, Kot W *et al.* A novel bacteriophage cocktail reduces and disperses *Pseudomonas aeruginosa* biofilms under static and flow conditions. *Microb Biotechnol* 2016; **9**: 61–74. https://doi.org/10.1111/1751-7915.12316
- Alemayehu D, Casey PG, McAuliffe O *et al.* Bacteriophages \$\phi MR299-2 and \$\phi NH-4\$ can eliminate *Pseudomonas aeruginosa* in the murine lung and on cystic fibrosis lung airway cells. *mBio* 2012; **3**: e00029-12. https://doi.org/10.1128/mBio.00029-12
- **115** Sakshi Bedi M, Verma V, Chhibber S. Amoxicillin and specific bacteriophage can be used together for eradication of biofilm of *Klebsiella pneumoniae* B5055. *World J Microbiol Technol* 2009; **25**: 1145–51. https://doi.org/10.1007/s11274-009-9991-8
- **116** Lu TK, Collins JJ. Dispersing biofilms with engineered enzymatic bacteriophage. *Proc Natl Acad Sci U S A* 2007; **104**: 11197–202. https://doi.org/10.1073/pnas.0704624104
- **117** Ryan EM, Alkawareek MY, Donnelly RF *et al.* Synergistic phage-antibiotic combinations for the control of *Escherichia coli* biofilms *in vitro. FEMS Immunol Med Microbiol* 2012; **65**: 395–8. https://doi.org/10.1111/j.1574-695X.2012.00977.x
- **118** Verma V, Harjai K, Chhibber S. Restricting ciprofloxacin-induced resistant variant formation in biofilm of Klebsiella pneumoniae B5055 by complementary bacteriophage treatment. *J Antimicrob Chemother* 2009; **64**: 1212–8. https://doi.org/10.1093/jac/dkp360
- **119** Coulter LB, McLean RJ, Rohde RE *et al.* Effect of bacteriophage infection in combination with tobramycin on the emergence of resistance in *Escherichia coli* and *Pseudomonas aeruginosa* biofilms. *Viruses* 2014; **6**: 3778–86. https://doi.org/10.3390/v6103778
- Zhang Y, Hu Z. Combined treatment of *Pseudomonas aeruginosa* biofilms with bacteriophages and chlorine. *Biotechnol Bioeng* 2013; **110**: 286–95. https://doi.org/10.1002/bit.24630
- Doub JB, Ng VY, Lee M *et al.* Salphage: salvage bacteriophage therapy for recalcitrant MRSA prosthetic joint infection. *Antibiotics (Basel)* 2022; **11**: 616. https://doi.org/10.3390/antibiotics11050616
- Tkhilaishvili T, Winkler T, Müller M *et al.* Bacteriophages as adjuvant to antibiotics for the treatment of periprosthetic joint infection caused by multidrug-resistant *Pseudomonas aeruginosa. Antimicrob Agents Chemother* 2019; **64**: e00924-19. https://doi.org/10.1128/AAC.00924-19
- Tkhilaishvili T, Wang L, Perka C *et al.* Using bacteriophages as a Trojan horse to the killing of dual-species biofilm formed by *Pseudomonas aeruginosa* and methicillin resistant *Staphylococcus aureus. Front Microbiol* 2020; **11**: 695. https://doi.org/10.3389/fmicb.2020.00695
- Meneses L, Brandão AC, Coenye T *et al.* A systematic review of the use of bacteriophages for *in vitro* biofilm control. *Eur J Clin Microbiol Infect Dis* 2023; **42**: 919–28. https://doi.org/10.1007/s10096-023-04638-1
- **125** Chegini Z, Khoshbayan A, Taati Moghadam M et al. Bacteriophage therapy against *Pseudomonas aeruginosa* biofilms: a review. *Ann Clin Microbiol Antimicrob* 2020; **19**: 45. https://doi.org/10.1186/s12941-020-00389-5
- Pearl S, Gabay C, Kishony R *et al.* Nongenetic individuality in the host-phage interaction. *PLoS Biol* 2008; **6**: e120. https://doi.org/10.1371/journal.pbio.0060120

- Dunsing V, Irmscher T, Barbirz S *et al.* Purely polysaccharide-based biofilm matrix provides size-selective diffusion barriers for nanoparticles and bacteriophages. *Biomacromolecules* 2019; **20**: 3842–54. https://doi.org/10.1021/acs.biomac.9b00938
- Vidakovic L, Singh PK, Hartmann R et al. Dynamic biofilm architecture confers individual and collective mechanisms of viral protection. *Nat Microbiol* 2018; **3**: 26–31. https://doi.org/10.1038/s41564-017-0050-1
- Chaudhry W, Lee E, Worthy A *et al.* Mucoidy, a general mechanism for maintaining lytic phage in populations of bacteria. *FEMS Microbiol Ecol* 2020; **96**: fiaa162. https://doi.org/10.1093/femsec/fiaa162
- Darch SE, Kragh KN, Abbott EA *et al.* Phage inhibit pathogen dissemination by targeting bacterial migrants in a chronic infection model. *mBio* 2017; **8**: e00240-17. https://doi.org/10.1128/mBio.00240-17
- Akturk E, Oliveira H, Santos SB *et al.* Synergistic action of phage and antibiotics: parameters to enhance the killing efficacy against mono and dual-species biofilms. *Antibiotics (Basel)* 2019; **8**: 103. https://doi.org/10.3390/antibiotics8030103
- Harada LK, Silva EC, Campos WF *et al.* Biotechnological applications of bacteriophages: state of the art. *Microbiol Res* 2018; **212-13**: 38–58. https://doi.org/10.1016/j.micres.2018.04.007
- Yehl K, Lemire S, Yang AC *et al.* Engineering phage host-range and suppressing bacterial resistance through phage tail fiber mutagenesis. *Cell* 2019; **179**: 459–469.e9. https://doi.org/10.1016/j.cell.2019.09.015
- Latka A, Lemire S, Grimon D *et al.* Engineering the modular receptorbinding proteins of *Klebsiella* phages switches their capsule serotype specificity. *mBio* 2021; **12**: e00455-21. https://doi.org/10.1128/mBio.00455-21
- Qin S, Liu Y, Chen Y et al. Engineered bacteriophages containing anti-CRISPR suppress infection of antibiotic-resistant *P. aeruginosa. Microbiol Spectr* 2022; **10**: e0160222. https://doi.org/10.1128/spectrum. 01602-22
- Mitsunaka S, Yamazaki K, Pramono AK *et al.* Synthetic engineering and biological containment of bacteriophages. *Proc Natl Acad Sci U S A* 2022; **119**: 62306730119. https://doi.org/10.1073/page.2306730119.
- 119: e2206739119. https://doi.org/10.1073/pnas.2206739119
- Motlagh AM, Bhattacharjee AS, Goel R. Biofilm control with natural and genetically-modified phages. *World J Microbiol Biotechnol* 2016; **32**: 67. https://doi.org/10.1007/s11274-016-2009-4
- Hagens S, Habel A, von Ahsen U *et al.* Therapy of experimental pseudomonas infections with a nonreplicating genetically modified phage. *Antimicrob Agents Chemother* 2004; **48**: 3817–22. https://doi.org/10.1128/AAC.48.10.3817-3822.2004
- Matsuda T, Freeman TA, Hilbert DW *et al.* Lysis-deficient bacteriophage therapy decreases endotoxin and inflammatory mediator release and improves survival in a murine peritonitis model. *Surgery* 2005; **137**: 639–46. https://doi.org/10.1016/j.surg.2005.02.012
- Peng H, Chen IA. Phage engineering and the evolutionary arms race. *Curr Opin Biotechnol* 2021; **68**: 23–9. https://doi.org/10.1016/j.copbio.2020.09.009
- Drulis-Kawa Z, Majkowska-Skrobek G, Maciejewska B. Bacteriophages and phage-derived proteins—application approaches. *Curr Med Chem* 2015; **22**: 1757–73. https://doi.org/10.2174/0929867322666150209152851
- Park JY, Moon BY, Park JW *et al.* Genetic engineering of a temperate phage-based delivery system for CRISPR/Cas9 antimicrobials against *Staphylococcus aureus. Sci Rep* 2017; **7**:44929. https://doi.org/10.1038/srep44929
- Cobb LH, Park J, Swanson EA *et al.* CRISPR-Cas9 modified bacteriophage for treatment of *Staphylococcus aureus* induced osteomyelitis and soft tissue infection. *PLoS One* 2019; **14**: e0220421. https://doi.org/10.1371/journal.pone.0220421
- Pei R, Lamas-Samanamud GR. Inhibition of biofilm formation by T7 bacteriophages producing quorum-quenching enzymes. *Appl Environ Microbiol* 2014; **80**: 5340–8. https://doi.org/10.1128/AEM.01434-14

- Pires DP, Monteiro R, Mil-Homens D *et al.* Designing *P. aeruginosa* synthetic phages with reduced genomes. *Sci Rep* 2021; **11**: 2164. https://doi.org/10.1038/s41598-021-81580-2
- Nick JA, Dedrick RM, Gray AL *et al.* Host and pathogen response to bacteriophage engineered against *Mycobacterium abscessus* lung infection. *Cell* 2022; **185**: 1860–1874.e12. https://doi.org/10.1016/j.cell.2022. 04.024
- **147** Emslander Q, Vogele K, Braun P *et al.* Cell-free production of personalized therapeutic phages targeting multidrug-resistant bacteria. *Cell Chem Biol* 2022; **29**: 1434–1445.e7. https://doi.org/10.1016/j.chembiol. 2022.06.003
- Liyanagedera SBW, Williams J, Wheatley JP *et al.* SpyPhage: a cell-free TXTL platform for rapid engineering of targeted phage therapies. *ACS Synth Biol* 2022; **11**: 3330–42. https://doi.org/10.1021/acssynbio. 2c00244
- Fa-Arun J, Huan YW, Darmon E *et al.* Tail-engineered phage P2 enables delivery of antimicrobials into multiple gut pathogens. *ACS Synth Biol* 2023; **12**: 596–607. https://doi.org/10.1021/acssynbio.2c00615
- Prokopczuk FI, Im H, Campos-Gomez J *et al.* Engineered superinfective Pf phage prevents dissemination of *Pseudomonas aeruginosa* in a mouse burn model. *mBio* 2023; **14**: e0047223. https://doi.org/10.1128/mbio.00472-23
- Balaban NQ, Helaine S, Lewis K *et al.* Definitions and guidelines for research on antibiotic persistence. *Nat Rev Microbiol* 2019; **17**: 441–8. https://doi.org/10.1038/s41579-019-0196-3
- Hall-Stoodley L, Costerton JW, Stoodley P. Bacterial biofilms: from the natural environment to infectious diseases. *Nat Rev Microbiol* 2004; **2**: 95–108. https://doi.org/10.1038/nrmicro821
- Akanda ZZ, Taha M, Abdelbary H. Current review—the rise of bacteriophage as a unique therapeutic platform in treating peri-prosthetic joint infections. *J Orthop Res* 2018; **36**: 1051–60. https://doi.org/10. 1002/jor.23755
- Chan WT, Espinosa M, Yeo CC. Keeping the wolves at bay: antitoxins of prokaryotic type II toxin-antitoxin systems. *Front Mol Biosci* 2016; **3**: 9. https://doi.org/10.3389/fmolb.2016.00009
- Song S, Wood TK. A primary physiological role of toxin/antitoxin systems is phage inhibition. *Front Microbiol* 2020; **11**: 1895. https://doi.org/10.3389/fmicb.2020.01895
- Castillo-Juárez I, Maeda T, Mandujano-Tinoco EA *et al.* Role of quorum sensing in bacterial infections. *World J Clin Cases* 2015; **3**: 575–98. https://doi.org/10.12998/wjcc.v3.i7.575
- Luthe T, Kever L, Thormann K *et al.* Bacterial multicellular behavior in antiviral defense. *Curr Opin Microbiol* 2023; **74**: 102314. https://doi.org/10.1016/j.mib.2023.102314
- Høyland-Kroghsbo NM, Maerkedahl RB, Svenningsen SL. A quorum-sensing-induced bacteriophage defense mechanism. *mBio* 2013; **4**: e00362-12. https://doi.org/10.1128/mBio.00362-12
- Saucedo-Mora MA, Castañeda-Tamez P, Cazares A *et al.* Selection of functional quorum sensing systems by lysogenic bacteriophages in *Pseudomonas aeruginosa. Front Microbiol* 2017; **8**: 1669. https://doi.org/10.3389/fmicb.2017.01669
- **160** Høyland-Kroghsbo NM, Bassler BL. Phage infection restores PQS signaling and enhances growth of a *Pseudomonas aeruginosa lasI* quorumsensing mutant. *J Bacteriol* 2022; **204**: e0055721. https://doi.org/10.1128/jb.00557-21

- Shah M, Taylor VL, Bona D *et al.* A phage-encoded anti-activator inhibits *quorum sensing* in *Pseudomonas aeruginosa. Mol Cell* 2021; **81**: 571–583.e6. https://doi.org/10.1016/j.molcel.2020.12.011
- Mion S, Rémy B, Plener L *et al.* Quorum quenching lactonase strengthens bacteriophage and antibiotic arsenal against *Pseudomonas aeruginosa* clinical isolates. *Front Microbiol* 2019; **10**: 2049. https://doi.org/10.3389/fmicb.2019.02049
- Qin X, Sun Q, Yang B *et al.* Quorum sensing influences phage infection efficiency via affecting cell population and physiological state. *J Basic Microbiol* 2017; **57**: 162–70. https://doi.org/10.1002/jobm. 201600510
- Severin GB, Ramliden MS, Ford KC *et al.* Activation of a *Vibrio cholerae* CBASS anti-phage system by quorum sensing and folate depletion. *mBio* 2023; **14**: e0087523. https://doi.org/10.1128/mbio.00875-23
- Xuan G, Lin H, Tan L *et al.* Quorum sensing promotes phage infection in *Pseudomonas aeruginosa* PAO1. *mBio* 2022; **13**: e0317421. https://doi.org/10.1128/mbio.03174-21
- Broniewski JM, Chisnall MAW, Høyland-Kroghsbo NM *et al.* The effect of quorum sensing inhibitors on the evolution of CRISPR-based phage immunity in Pseudomonas aeruginosa. *ISME J* 2021; **15**: 2465–73. https://doi.org/10.1038/s41396-021-00946-6
- Ghosh D, Roy K, Williamson KE *et al.* Acyl-homoserine lactones can induce virus production in lysogenic bacteria: an alternative paradigm for prophage induction. *Appl Environ Microbiol* 2009; **75**: 7142–52. https://doi.org/10.1128/AEM.00950-09
- Hu J, Ye H, Wang S *et al.* Prophage activation in the intestine: insights into functions and possible applications. *Front Microbiol* 2021; **12**: 785634. https://doi.org/10.3389/fmicb.2021.785634
- **169** Rossmann FS, Racek T, Wobser D *et al.* Phage-mediated dispersal of biofilm and distribution of bacterial virulence genes is induced by quorum sensing. *PLoS Pathog* 2015; **11**: e1004653. https://doi.org/10.1371/journal.ppat.1004653
- Fernández-Piñar R, Cámara M, Dubern JF et al. The Pseudomonas aeruginosa quinolone quorum sensing signal alters the multicellular behaviour of Pseudomonas putida KT2440. Res Microbiol 2011; **162**: 773–81. https://doi.org/10.1016/j.resmic.2011.06.013
- Silpe JE, Bassler BL. A host-produced quorum-sensing autoinducer controls a phage lysis-lysogeny decision. *Cell* 2019; **176**: 268–280.e13. https://doi.org/10.1016/j.cell.2018.10.059
- Maffei E, Woischnig A-K, Burkolter MR *et al.* Phage Paride can kill dormant, antibiotic-tolerant cells of *Pseudomonas aeruginosa* by direct lytic replication. *Nat Commun* 2024; **15**: 175. https://doi.org/10.1038/s41467-023-44157-3
- Pacios O, Fernández-García L, Bleriot I *et al.* Enhanced antibacterial activity of repurposed mitomycin C and imipenem in combination with the lytic phage vB\_KpnM-VAC13 against clinical isolates of *Klebsiella pneumoniae*. *Antimicrob Agents Chemother* 2021; **65**: e0090021. https://doi.org/10.1128/AAC.00900-21
- Briers Y, Walmagh M, Grymonprez B *et al.* Art-175 is a highly efficient antibacterial against multidrug-resistant strains and persisters of *Pseudomonas aeruginosa. Antimicrob Agents Chemother* 2014; **58**: 3774–84. https://doi.org/10.1128/AAC.02668-14
- Schmelcher M, Donovan DM, Loessner MJ. Bacteriophage endolysins as novel antimicrobials. *Future Microbiol* 2012; **7**: 1147–71. https://doi.org/10.2217/fmb.12.97