Title Nove Research Theme IIAR Project Type Wet	IIAR26Ex Westra el phage-based therapeutics against drug-resistant bacteria
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Summary Anti	lab
thre project that resist effect use sense general chall approximations.	piotic-resistant Pseudomonas aeruginosa is a major global health at, causing infections that are increasingly difficult to treat. This ect will develop next-generation bacteriophage (phage) therapies target and kill these bacteria, overcoming conventional antibiotic tance. You will explore how phages can be engineered to maximise ctiveness, understand how bacteria evolve resistance to phages, and synthetic biology to design precision tools that restore phage itivity. This interdisciplinary research integrates microbiology, emics, and bioengineering to tackle one of the most urgent enges in infectious disease, offering training in cutting-edge oaches with clear real-world impact.
a cri bact hosp antil gain signi bact mod can Ther to or phag host mod path ende synt platt xend epig cons This mod build deve engi CRIS that robu bypa Rese Can	microbial resistance (AMR) is an urgent global health threat, driving tical need for innovative therapeutics to control multidrug-resistant erial pathogens. Pseudomonas aeruginosa is a major cause of ital-acquired infections and is inherently resistant to many piotics. As conventional treatments fail, bacteriophages (phages) are ng renewed attention as precision antimicrobials. However, a ficant barrier to effective phage therapy is the remarkable ability of eria to deploy defence systems, such as CRISPR—Cas, restriction—iffication, and abortive infection, to neutralise phages before they clear infection. e is an increasing interest in the use of engineered phage genomes vercome these defences. Yet scaling up production of modified tes remains challenging. Traditional in vivo production in bacterial is imposes limitations, including toxicity of certain genetic iffications (e.g., broad-range methyltransferases), the need to use ogenic strains or species as production platforms, and the release of otoxins by the production host. To address these obstacles, cell-free netic biology systems have emerged as a powerful alternative. These orms enable assembly, replication and rebooting of non-cognate uphages entirely in vitro, allowing precise genome editing and enetic modification in a controlled environment without the traints of bacterial viability. project will pioneer an E. coli-based cell-free system to generate iffied P. aeruginosa phages with enhanced therapeutic properties, ling on advanced phage production workflows and cell-free systems eloped by Dr. Temperton. The approach will combine genome neering (e.g., adding anti-defence genes or payloads like toxins or PR-silencing) with epigenomic tailoring (e.g., methylation patterns evade host restriction). Ultimately, this work aims to produce a st pipeline for designing and manufacturing bespoke phages that is bacterial defences and selectively kill target pathogens. arch Question: we create a scalable, programmable cell-free platform for producing neered P.

modifications to overcome bacterial defences and improve therapeutic efficacy?

Specific Objectives:

Objective 1: Establish an E. coli cell-free phage production system Addition of non-cognate sigma factors to cell free systems can reprogram E. coli RNA polymerase to recognise xenophage promoters in order to initiate rebooting. Here, the student will adapt and optimise existing E. coli cell-free lysates to reboot P. aeruginosa phages via supplementation with plasmids encoding both sigma factors native to Ps. aeruginosa and novel sigma factors designed through generative AI (e.g. evo2).

Objective 2: Engineer phage genomes to encode counter-defence systems and antimicrobial payloads

Using synthetic biology and in vitro recombination techniques, the student will generate phage genomes incorporating genes that neutralise bacterial defences (e.g., anti-CRISPR proteins) or deliver toxins to enhance bacterial clearance.

Objective 3: Tailor phage epigenomes to bypass restriction-modification defences

The project will explore methods to methylate phage DNA with promiscuous methyltransferases in vitro, building on prior attempts to do this in vivo. This will involve characterising methylation patterns, testing how they impact the efficacy of rebooting, and whether phage modification improves treatment efficacy against resistant P. aeruginosa strains

Objective 4: Validate engineered phages against clinically relevant P. aeruginosa strains

The student will evaluate the infectivity, replication kinetics, and bactericidal activity of modified phages in vitro against an international panel of clinical isolates of P. aeruginosa, comparing their performance to unmodified controls. These studies will clarify the contribution of epigenetic and genetic modifications to overcoming bacterial defences. Student Ownership:

This project offers multiple avenues for the student to take initiative and shape the work:

Developing new cell-free reaction protocols and optimising production workflows.

Designing novel genetic constructs and testing different anti-defence combinations.

Exploring the impact of specific methylation patterns on phage—host interactions.

Contributing to publications, protocols, and intellectual property emerging from the work.

Collaborative Environment and Training:

The student will be supervised by Westra, Temperton and Szczelkun. We foster a positive research culture that prioritises mutual respect, collegiality, and the highest standards of research integrity. We are dedicated to creating a supportive environment where everyone feels valued and empowered to contribute, and actively promote equality, diversity, and inclusion in recruitment, supervision, and daily practice, recognising that diverse perspectives drive scientific excellence. Our

collaborative approach to team science encourages interdisciplinary exchange, shared ownership of ideas, and collective success. Through this collaboration, the student will gain cutting-edge skills in: Cell-free synthetic biology and phage manufacturing. Genetic and epigenetic engineering of phage genomes. Microbiological assays and infection modelling.

Genomic and epigenomic characterisation of phage particles.

The student will be embedded in an exceptional network of research activity and expertise through:

- 1) the BBSRC-funded sLoLa "MultiDefence" project, led by Prof. Westra, which investigates the mechanisms and evolution of the P. aeruginosa immune system against phage infections (https://sites.exeter.ac.uk/multidefence/).
- 2) the Safephage project (led by Mike Brockhurst, Westra co-I), which focuses on the safe and effective development of synthetic phage therapies using a yeast-based rebooting platform and in vivo phage production host.
- 3) The Citizen Phage Library project (led by Ben Temperton), which develops bespoke Pseudomonas phage treatments.

Supervisory Team	
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