Industry-Academia Structural Biology Alliance to Tackle Bacterial Antibiotic Resistance

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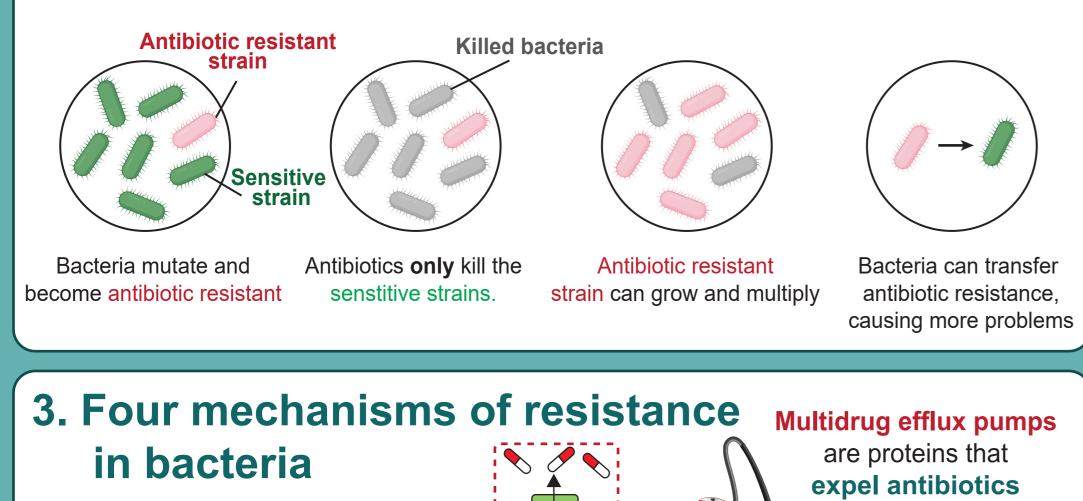
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1. What is antimicrobial resistance?

Antimicrobial resistance (AMR) occurs when microbes such as bacteria and viruses accumulate changes over time. They no longer respond to antimicrobial treatments such as antibiotics and antivirals designed to kill them. Main causes include the misuse of antimicrobials in humans, animals and plants.

Antibiotic resistance in bacteria

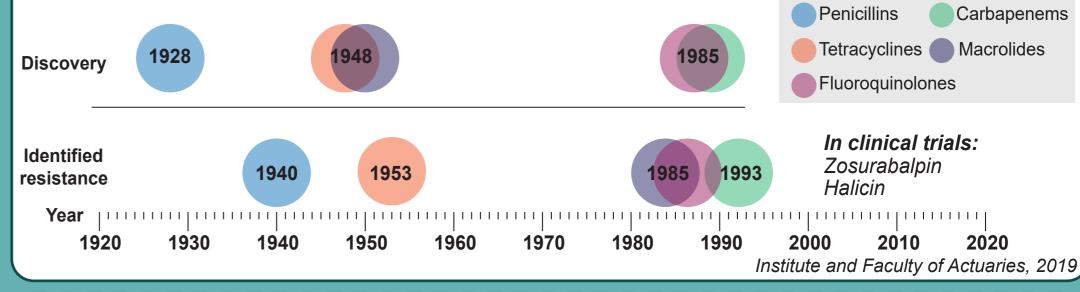


2. Why is it a global challenge?

- AMR is one of the top ten threats to global public health, food security and development. It is known as a "silent pandemic" (WHO).
- In 2019, bacterial antibiotic resistance was directly responsible for 1.27 million global deaths.
- Antibiotic resistance makes medical procedures and treatments such as surgery and cancer chemotherapy riskier.

Antibiotic discovery and resistance timeline

- Bacteria are getting faster at developing resistance to antibiotics.
- 30 years since a new class of antibiotics was last introduced.

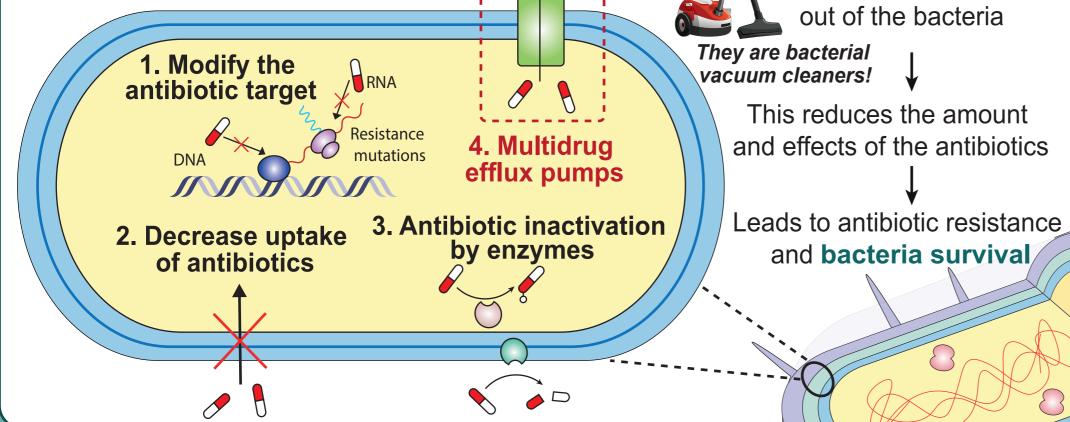


WORLDWIDE

10 million

deaths per year by 2050 (greater than cancer)

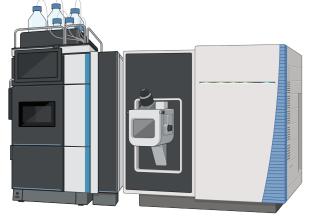
Costing £66 trillion (www.gov.uk)



5. Structural biology methods

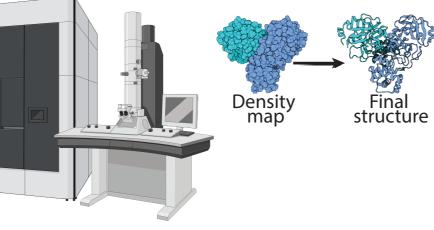
Structural biology is a key tool in **drug discovery** which looks at proteins at a molecular level, unlocking secrets of structure, function and interactions.

Hydrogen/Deuterium eXchange Mass Spectrometry (HDX-MS)



Tool for analysing structural features and dynamic (movement) properties of proteins

Cryogenic-electron microscopy (cryo-EM)



Super microscope to visualise how atoms of a protein and a drug are arranged in 3D space

What is the structure of MdtF?

Same class of protein as AcrB, but currently no structural information is available.

4. Aims

How do **multidrug efflux pumps** work to expel antibiotics?

Can we **inhibit (stop)** that happening to allow antibiotics to remain inside and kill bacteria?

To tackle this biological problem we:

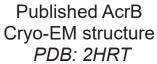
- Establish a multi-disciplinary academia /industry partnership
- Focus on two multidrug efflux pumps: AcrB and MdtF
 - Use state-of-the-art structural biology methods

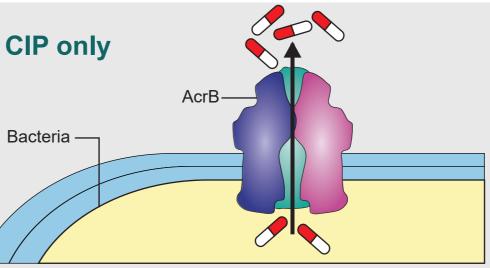
6. Research findings How does <u>AcrB</u> expel antibiotics?

Although structures exist for AcrB, its movement and function are unknown.

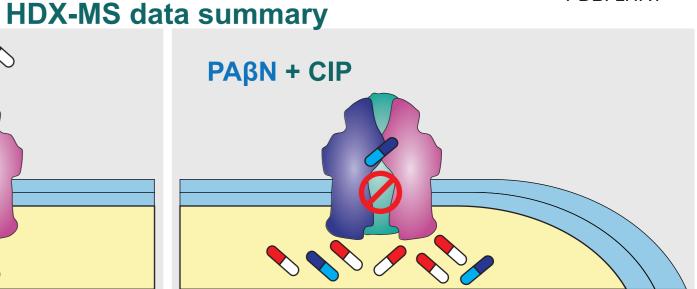
• We used HDX-MS to explore how the antibiotic (CIP) and the efflux pump inhibitor (PABN) affect AcrB movements.





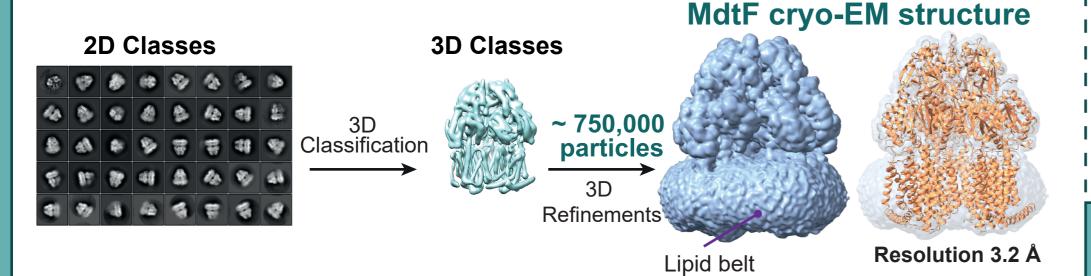


AcrB movement increases Antibiotic **CIP** is expelled out of the bacteria Lower cell concentration of CIP Bacteria is antibiotic resistant and survives



PAβN restricts AcrB movement Antibiotic CIP remains inside the bacteria CIP can perform its role in killing bacteria Drug efflux is inhibited and bacteria dies

• Increased expression and efflux in anaerobic/acidic conditions (e.g. mammalian gut).



- We solved the first structure of MdtF using cryo-EM.
- Cryo-EM structure reveals three monomers (similar to AcrB).

Future work

nspired by Patients

Driven by **Science**

- Perform HDX-MS to investigate movement and function.
- Find inhibitors that restrict the movement of MdtF and stop the removal of antibiotics, consequently prevent bacterial resistance.

University of Southampton

Antibiotic susceptibility assays: PAβN enhances CIP activity

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Microbial	Minimum Inhibitory
strain	Concentration (µg/ml)
	CIP
E. coli ATCC 25922	0.008
_ / + 16 μg/ml	0.004
Aggradue (19 μg/m) + 32 " + 64 "	0.002
	0.001
^۳ ق +128 "	≤ 0.00012

Impact

- Efflux pumps are also found in human cells (not just in bacteria).
- The leading cause of cancer chemotherapy failure is the development of multidrug resistance by efflux pumps e.g. P-glycoproteins in human cancer cells.
- Overall, understanding how inhibitors of efflux pumps work can:
 - overcome bacterial resistance and allow us to reuse antibiotics.
 - restore the sensitivity of human cancer cells toward chemotherapy drugs.

References



Antibiotic Resistance: Modelling the Impact on Mortality and Morbidity A report by the Antibiotic Resistance Working Party. Institute and faculty of Actuaries, 2019. Antimicrobial Resistance Collaborators. (2022). Global burden of bacterial antimicrobial resistance in 2019: a systematic analysis. The Lancet; 399(10325): P629-655. Reading, E., Ahdash, Z., et al. Perturbed structural dynamics underlie inhibition and altered efflux of the multidrug resistance pump AcrB. Nat Commun 11, 5565 (2020).

The efflux pump inhibitor PAβN can stop AcrB from expelling the antibiotic CIP, preventing bacterial resistance.

Reading, E, Ahdash, Z et al. Nature Commun, 2020