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# Evolutionary principles of antimicrobial peptide resistance

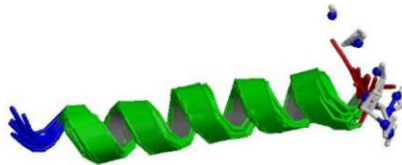
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# Antimicrobial peptides (AMP)

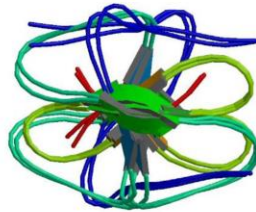
(A)  $\alpha$ -helical



Magainin-2

DOI:10.2210/pdb2mag/pdb

(B)  $\beta$ -sheet



$\beta$ -defensin 1

DOI:10.2210/pdb1kj5/pdb

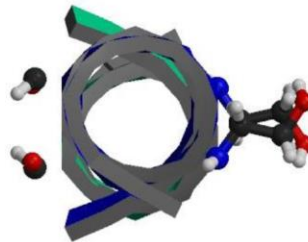
(C) extended



Indolicidin

DOI:10.2210/pdb1g89/pdb

(D) loop



Gramicidin

DOI:10.2210/pdb1mag/pdb

Short and diverse

Broad spectrum of  
bactericidal activity

Found in bacteria and  
eukaryotes alike

Part of innate immune  
system in vertebrates

# Some used in the clinic

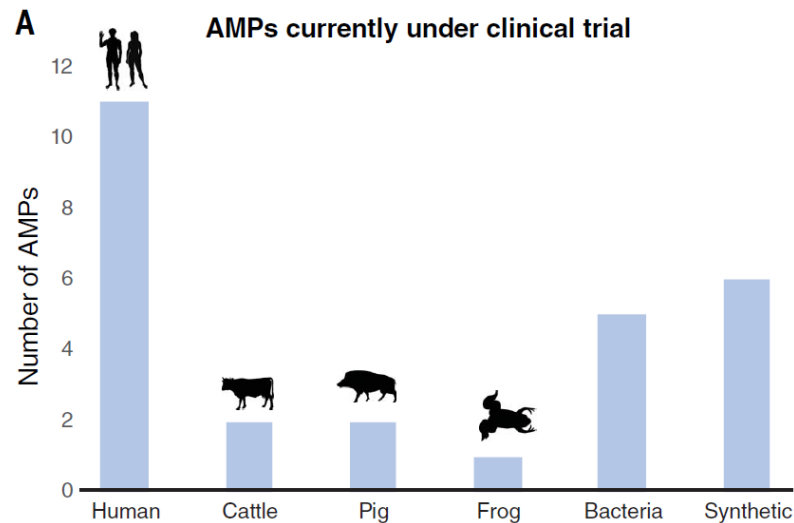


A last-resort antibiotics against multi-drug resistant Gram-negatives

# Main concerns

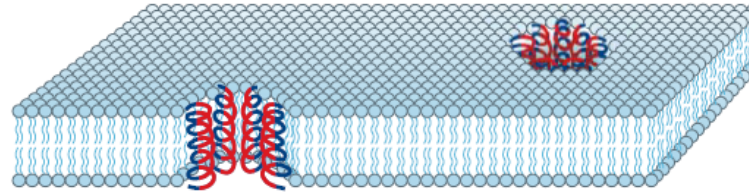
1) Resistance to certain AMPs evolves in the laboratory and in nature

2) Resistance may compromise natural immunity (i.e. cross-resistance to human peptides)

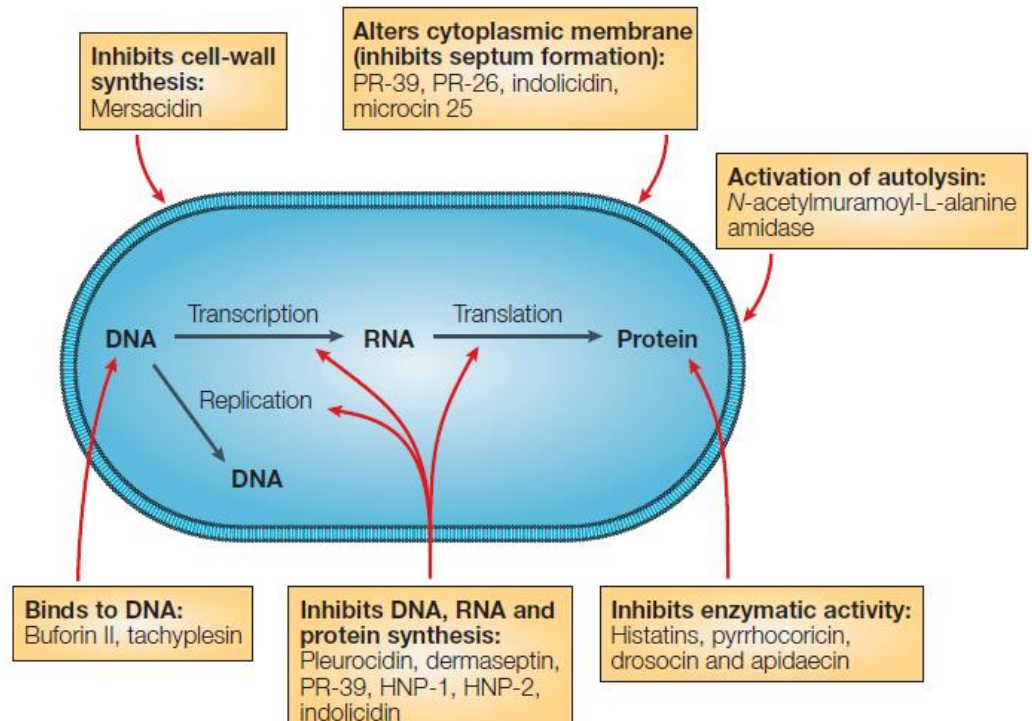


# AMPs have diverse modes of action

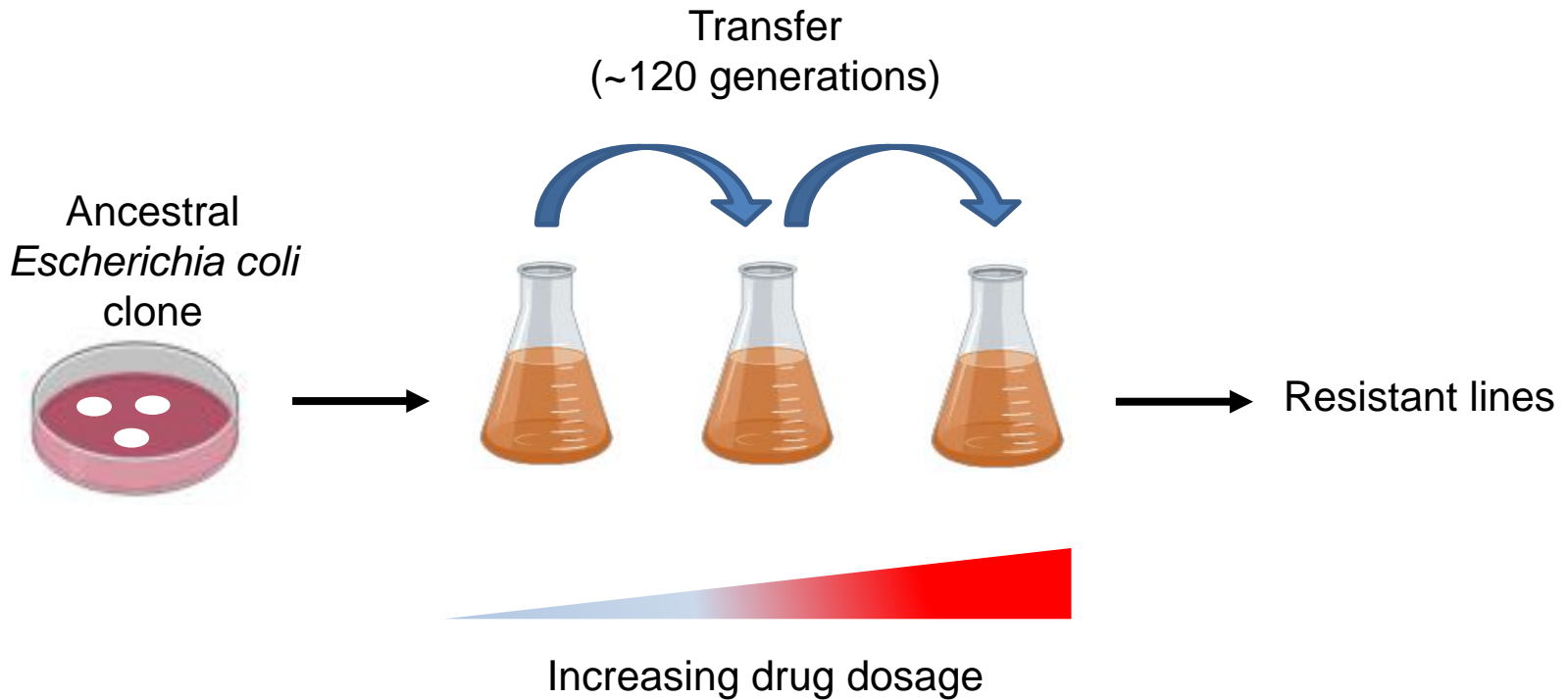
## Membrane targeting / pore formers



## Intracellular targeting



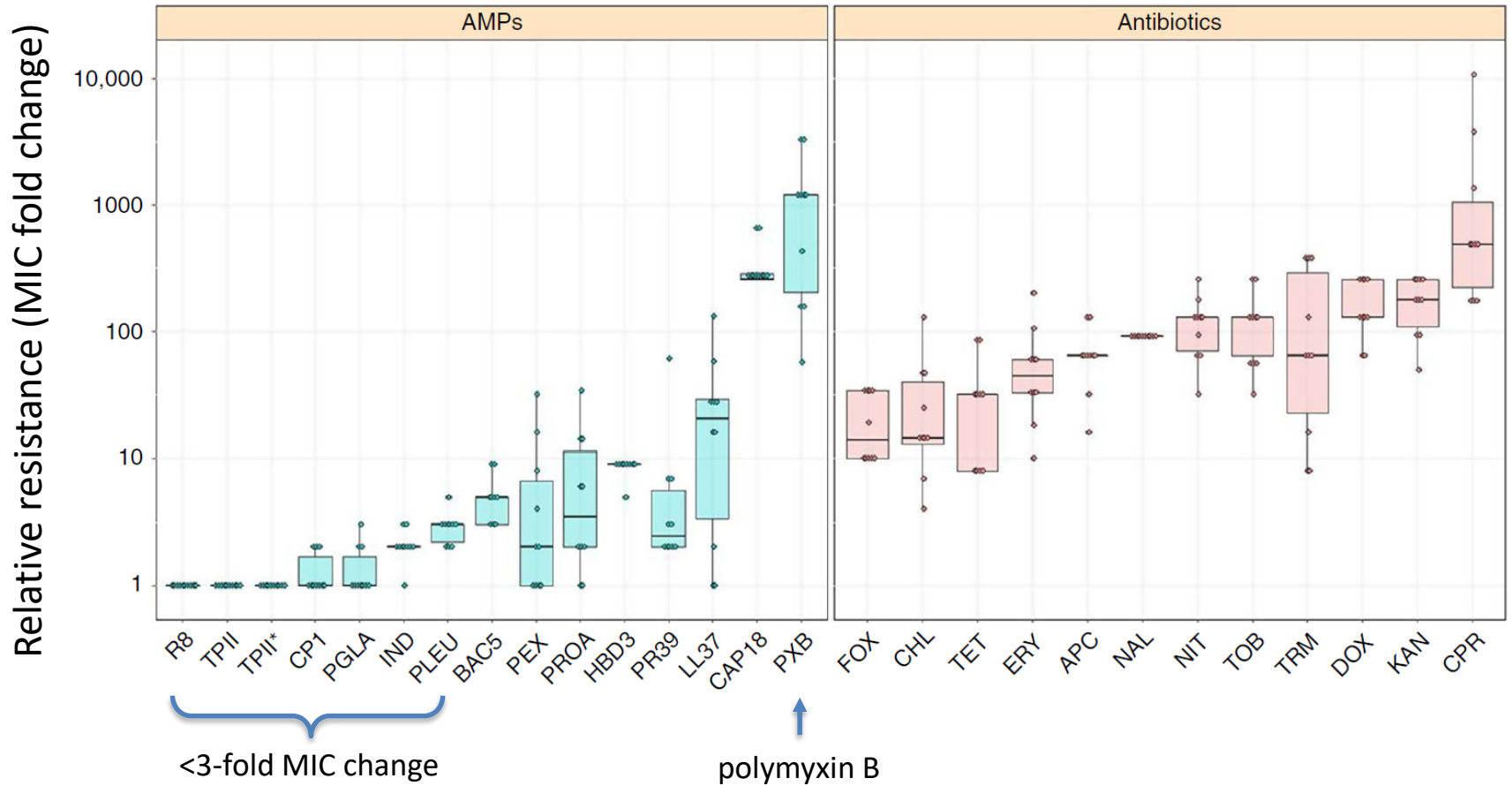
# Laboratory evolution of resistance



10 parallel lines per drug

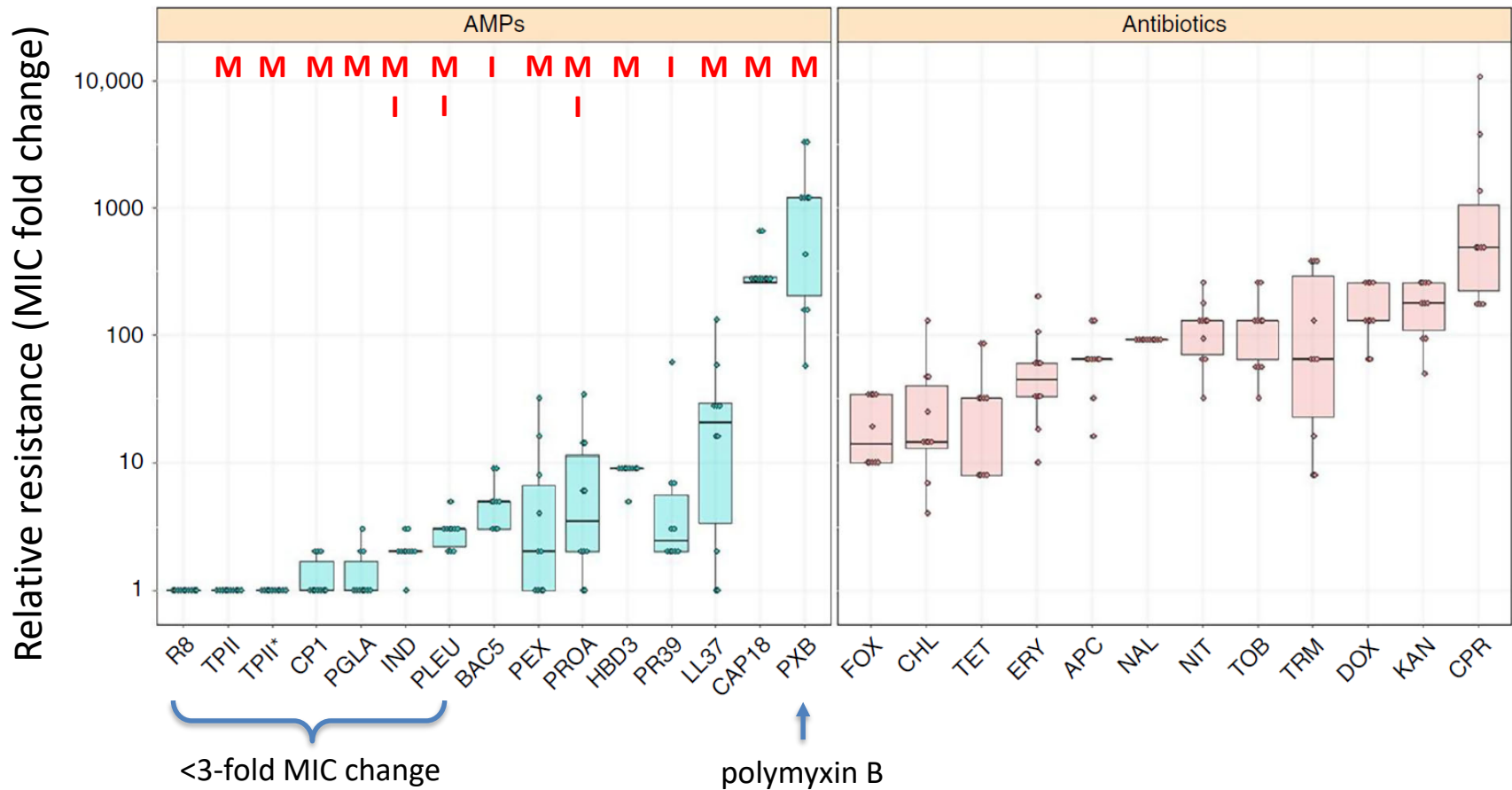
14 AMPs with diverse structures and mechanisms

# Lower resistance potential for AMPs than for antibiotics



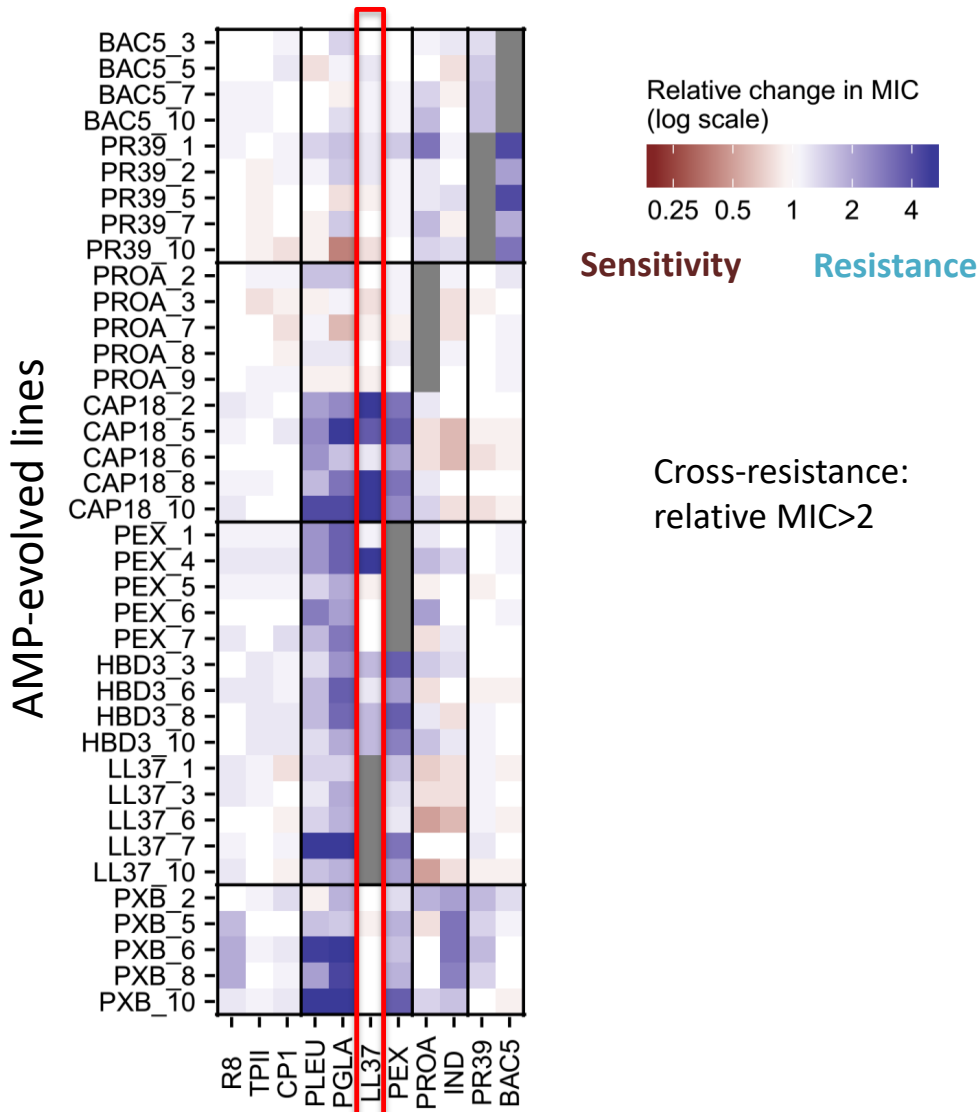
# Membrane targeting AMPs are not universally resistance-free

**M: membrane targeting, I: intracellular targeting**





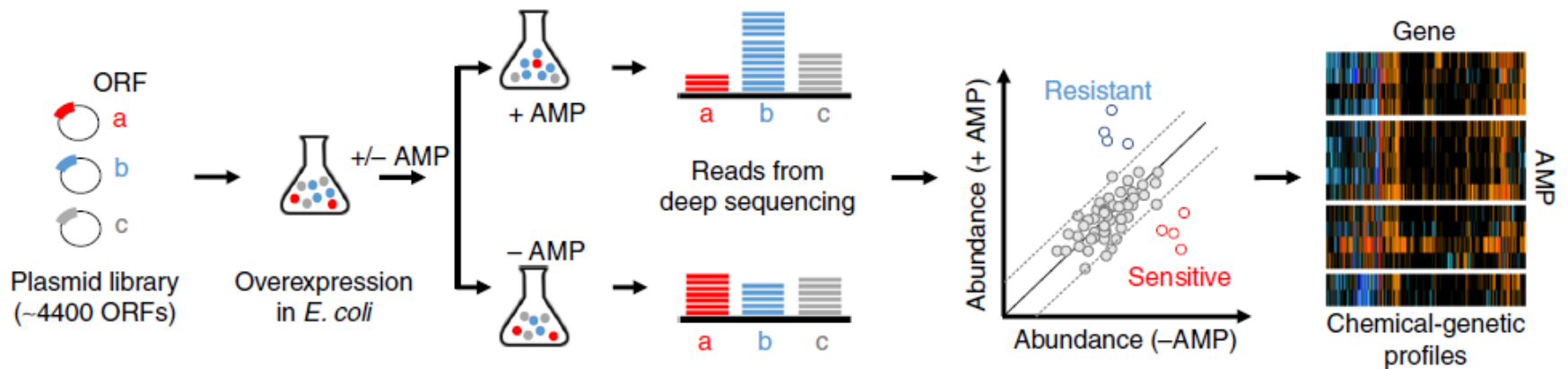
# Cross-resistance is relatively rare



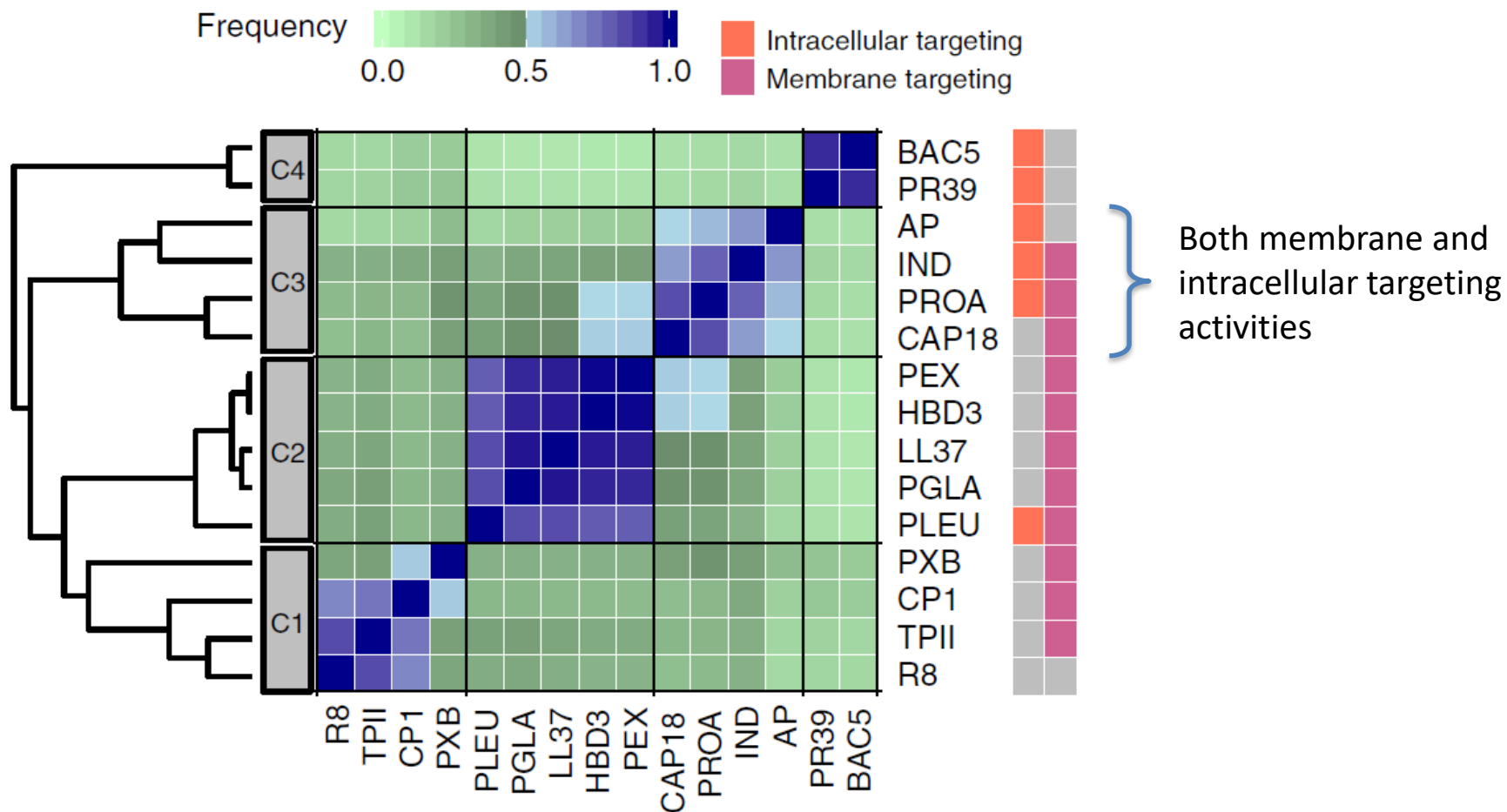
Only 16.5% of combinations show cross-resistance

But human peptide LL37 shows cross-resistance to several lines

# Mapping resistance-modulating genes through chemical genetics

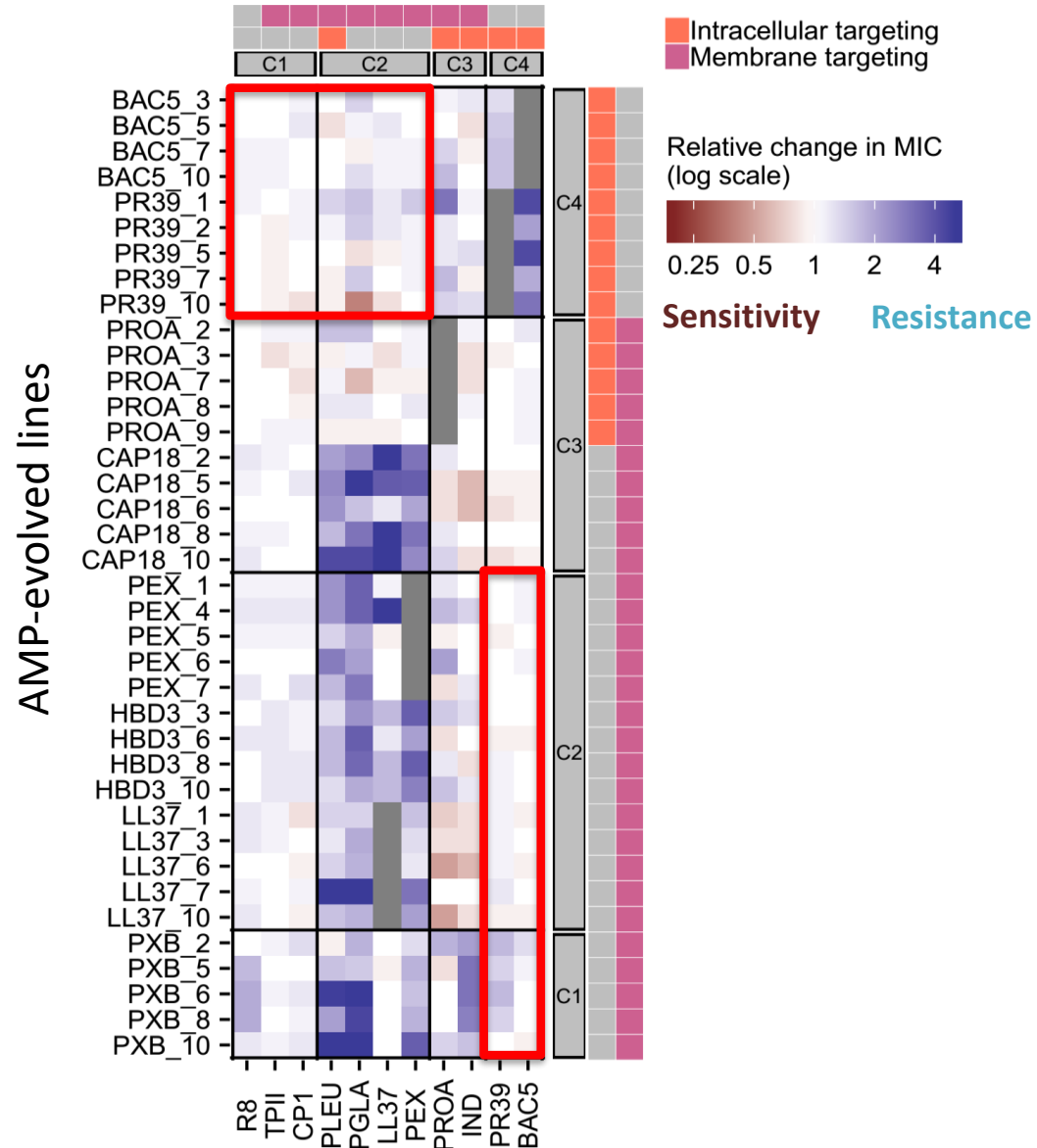
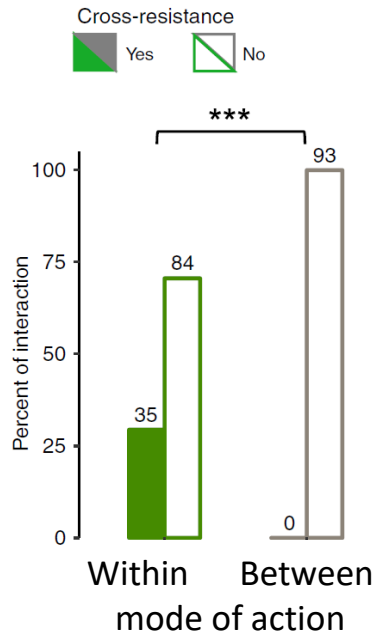


# Chemical-genetic profiles group AMPs with similar mode of action...

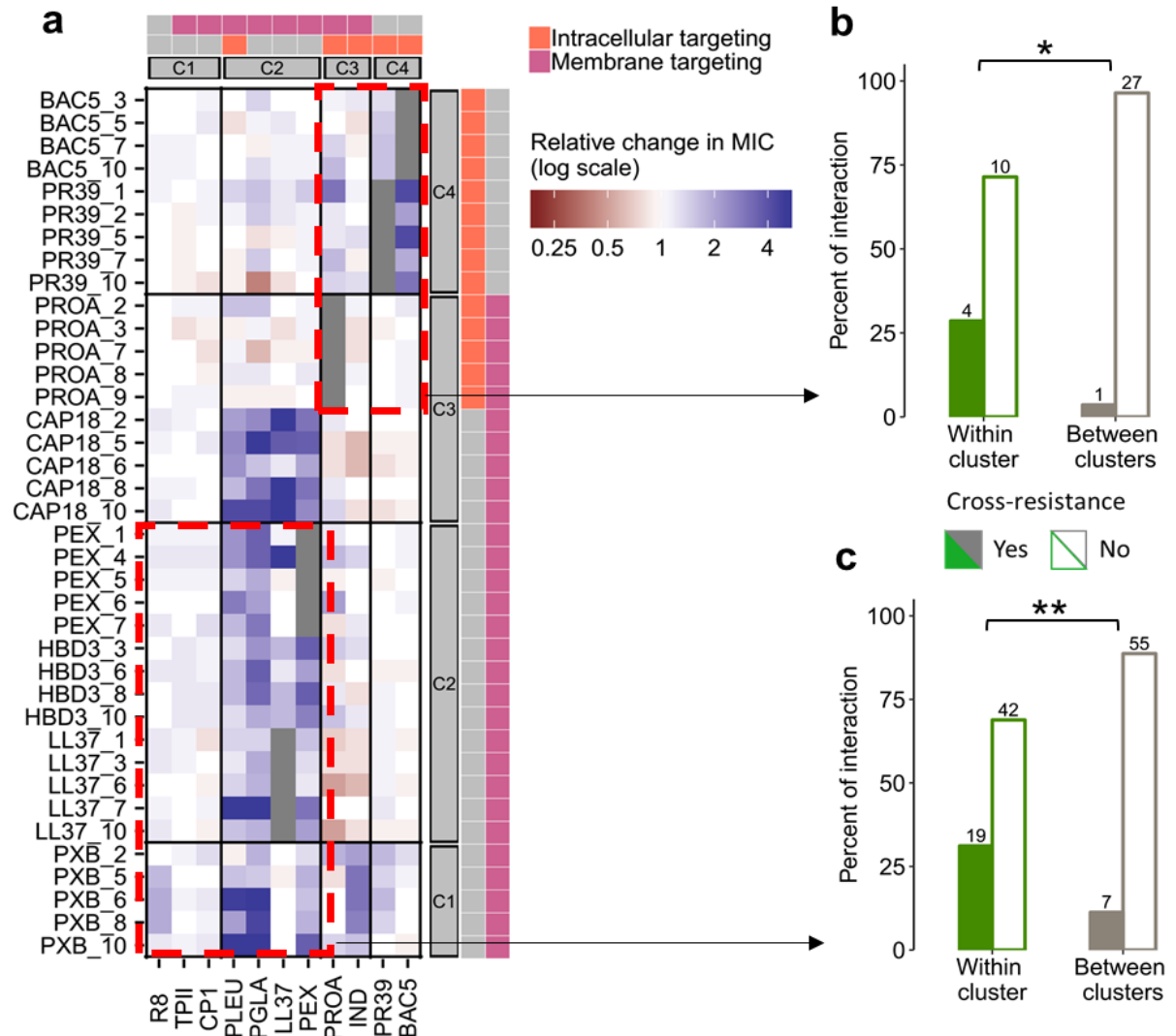


# ...and informs on cross-resistance

1) No cross-resistance between strictly intracellular and membrane targeting AMPs



## 2) Chemical-genetic profiles also predicts cross-resistance patterns within mode of action



# Conclusions

- 1) AMPs differ greatly in resistance propensity
- 2) Cross-resistance to human peptides could be minimized by deploying intracellular targeting AMPs

# Acknowledgements

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wellcome trust







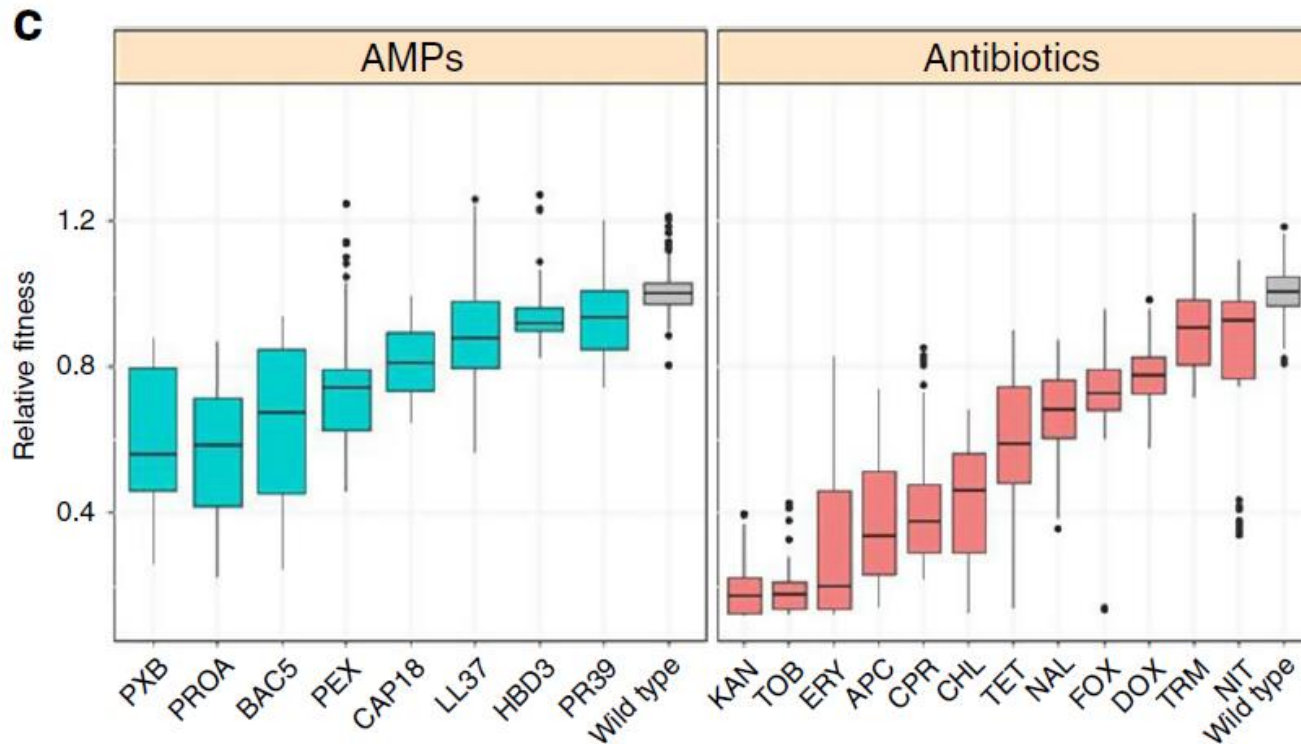
**Table 1 List and characteristics of AMPs used in this study. Their abbreviation, described mode of action, and clinical relevance (for details see Supplementary Data 7).**

<b>Name of AMP</b>	<b>Abbreviation</b>	<b>Mode of action</b>	<b>Clinical relevance</b>
Apidaecin IB	AP	Inhibits protein biosynthesis by targeting ribosomes; Interacts with DnaK, GroEL/ GroES, FtsH	Yes
Bactenecin 5	BAC5	Inhibits protein and RNA synthesis	n.a.
CAP18	CAP18	Disrupts cell membrane	Yes
Cecropin P1	CP1	Disrupts cell membrane	n.a.
Human beta-defensin-3	HBD-3	Disrupts cell membrane; Inhibits lipid II in peptidoglycan biosynthesis	n.a.
Indolicidin	IND	Inhibits DNA and protein synthesis; Disrupts cell membrane; Inhibits septum formation	Yes
LL-37 human cathelicidin	LL37	Disrupts cell membrane; Induces ROS formation	Yes
Peptide glycine-leucine amide	PGLA	Disrupts cell membrane	n.a.
Pexiganan	PEX	Disrupts cell membrane	Yes
Pleurocidin	PLEU	Disrupts cell membrane; Induces ROS formation; Inhibits protein and DNA synthesis	n.a.
Polymyxin B	PXB	Disrupts cell membrane; Induces ROS formation	Yes
PR-39	PR39	Inhibits protein and DNA synthesis	n.a.
Protamine	PROA	Affects cellular respiration and glycolysis; Disrupts cell envelop	n.a.
R8	R8	n.a.	n.a.
Tachyplesin II	TPII	Disrupts cell membrane	n.a.

*n.a.* no data available

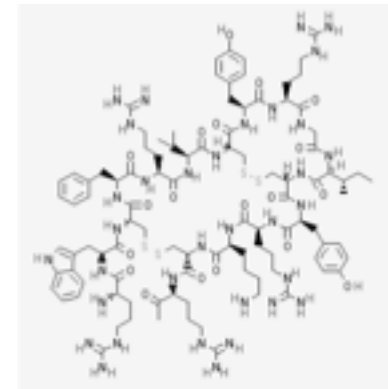
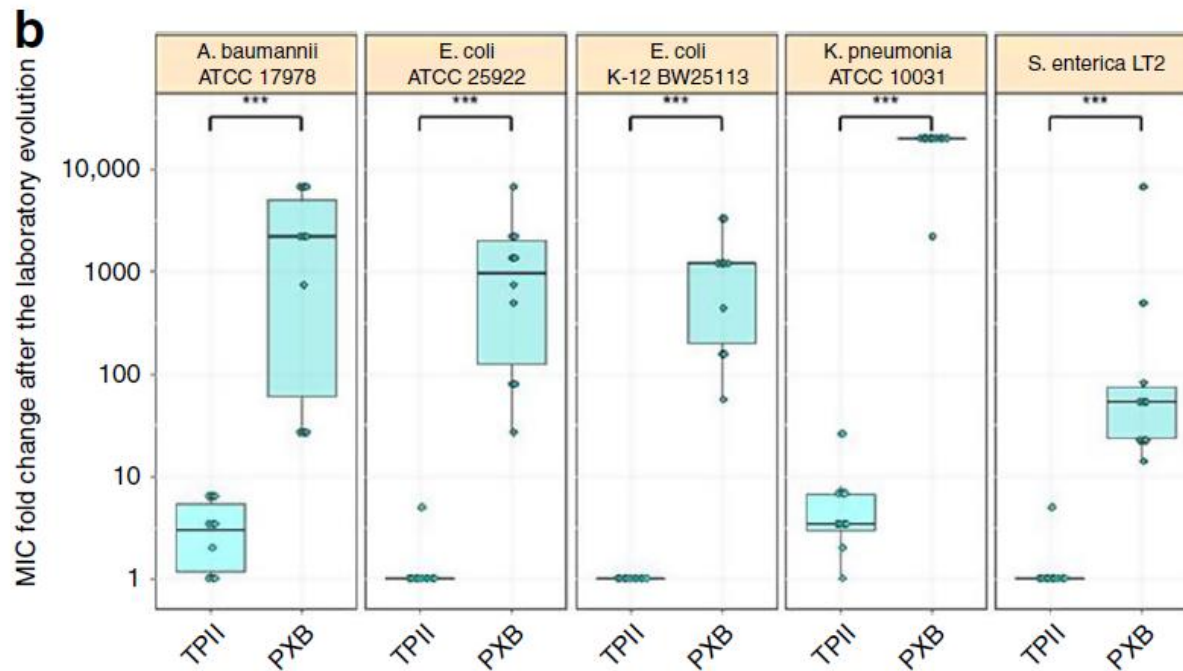
# Lower resistance potential of AMPs is not explained by high cost of resistance

In general, AMP evolved lines show lower fitness costs



# Tachyplesin II: a cationic $\beta$ -hairpin antimicrobial peptide discovered in horseshoe crab

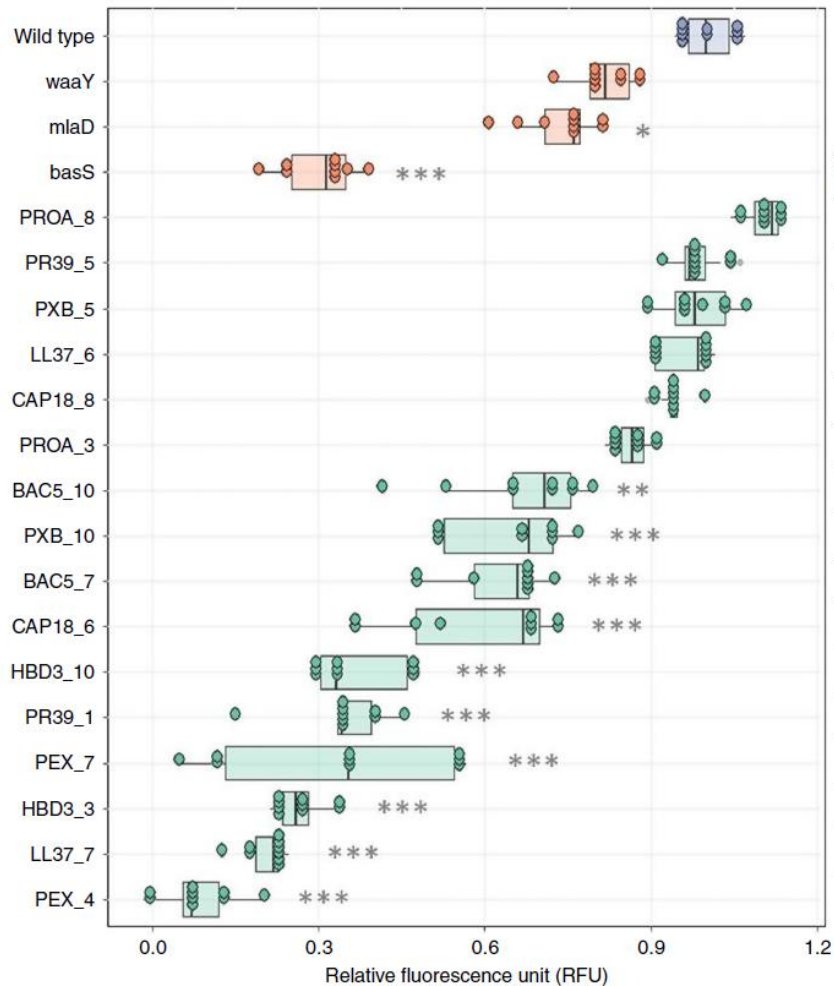
Appears resistance-free when evolution is conducted in in different clinical isolates



Tachyplesin II

# Surface charge changes underlie resistance

- 13 out of 16 tested evolved lines show a reduced net negative surface charge
- Carry mutations in BasR-BasS two-component system, a regulator of LPS pathway / phospholipid trafficking

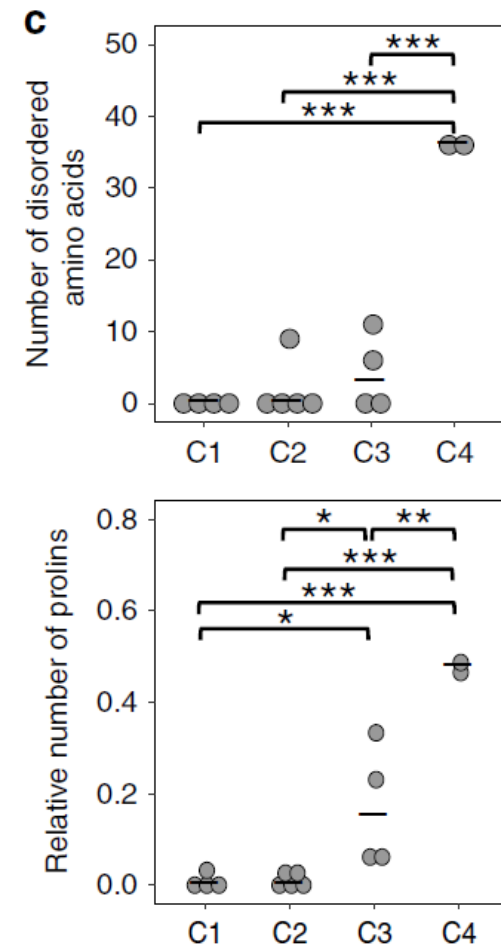
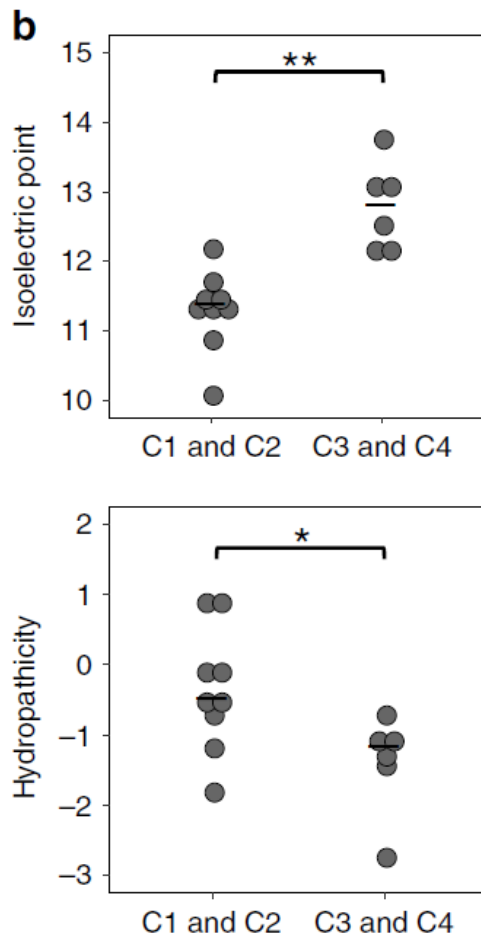


waa genes			mla genes				bas genes	
waaQ	waaY	waaJ	mlaE	mlaF	mlaD	mlaA	basS	basR
■							■	
	■		■					
■		■						
	■							
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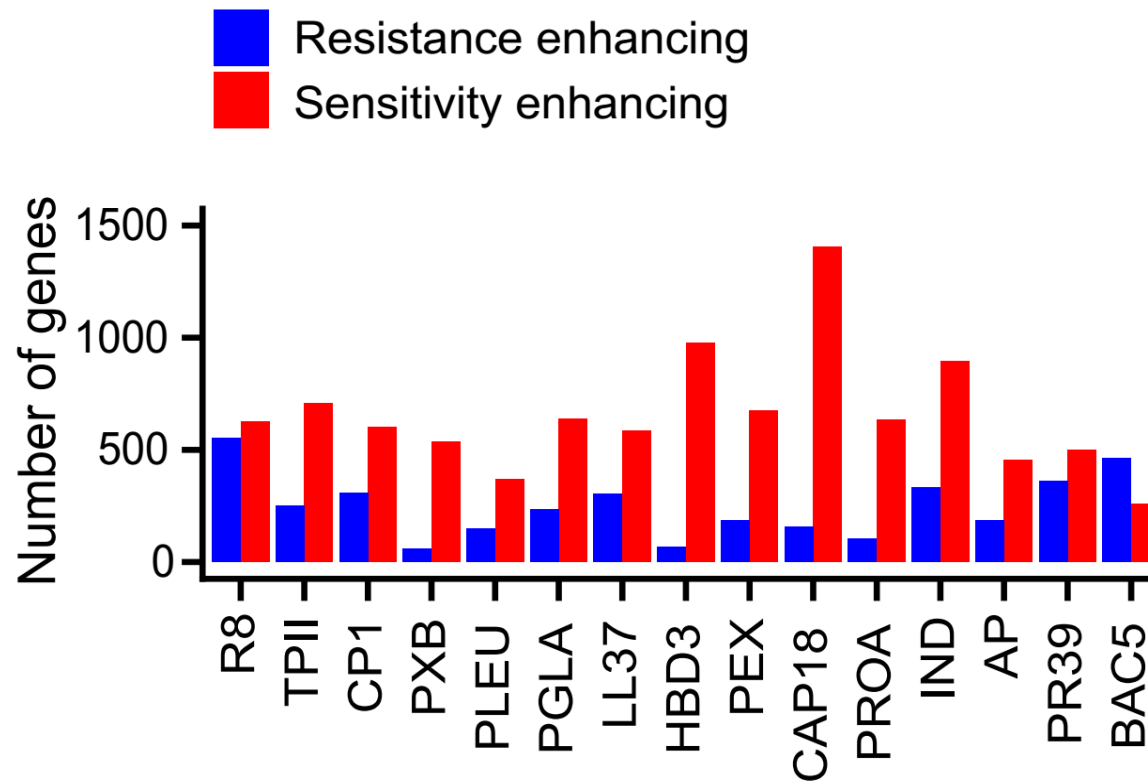
# Chemical-genetic profiling distinguish AMPs with different physicochemical properties

C1 + C2 = membrane targeting

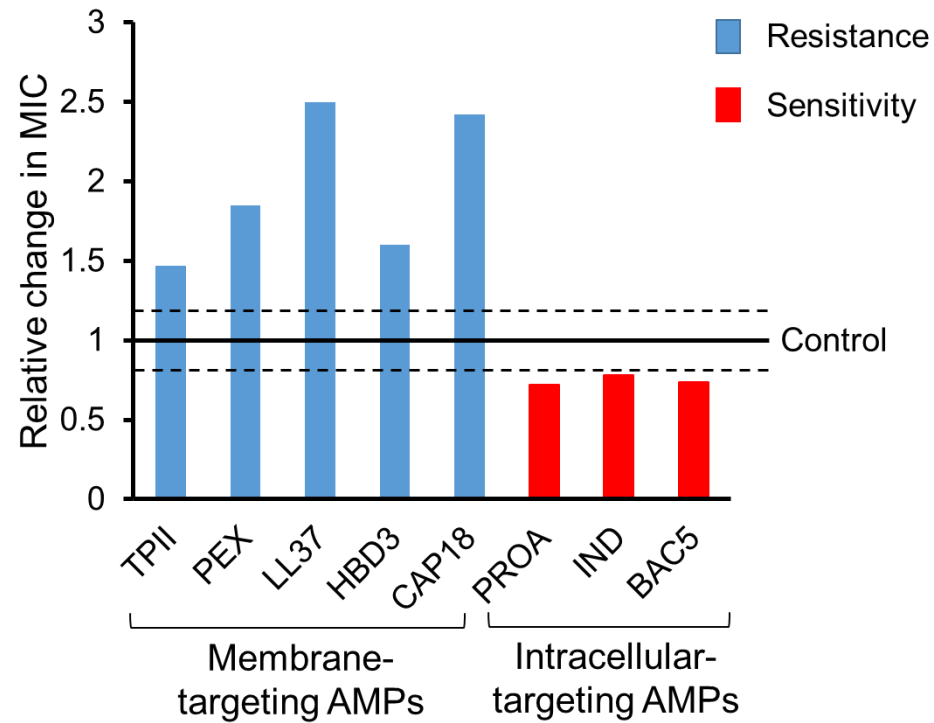
C3 and C4 intracellular targeting AMPs differ: C4 is more prone to disorder



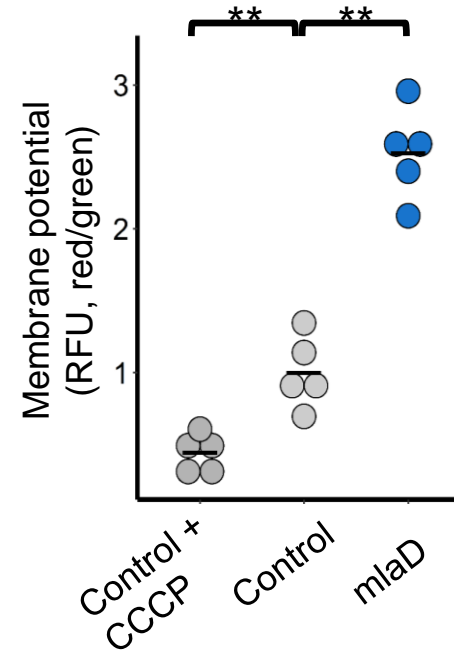
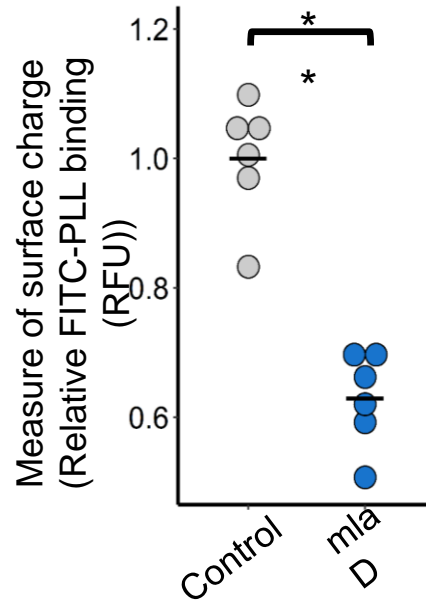
# A large set of genes modulate AMP susceptibility



# Perturbation of *miaD* confers resistance to membrane-targeting AMPs and sensitivity to intracellular-targeting AMPs

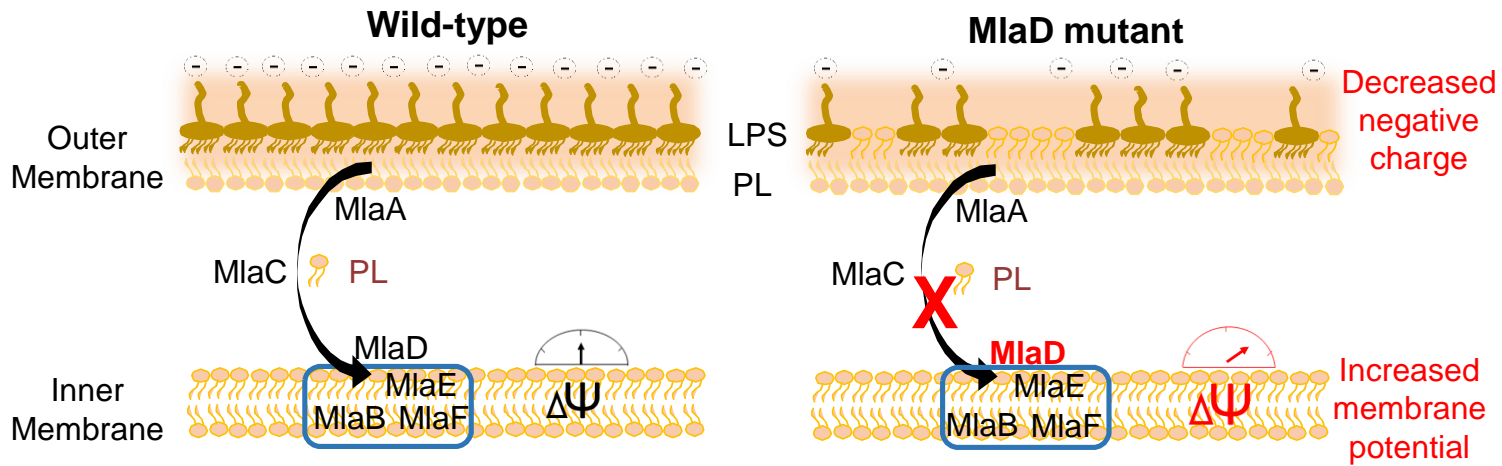


# Perturbation of *miaD* decreases membrane surface charge and increases membrane potential





# Molecular basis of the collateral sensitivity interaction



# Functionally diverse AMPs and antibiotics used

AMPs		Antibiotics	
Polymyxin B	(cyclic bacterial)	Trimetoprin	(FolA inhibitor)
PGLa	(pore former, amphibian)	Ampicillin	(beta lactam)
LL37	(pore former, human)	Tetracycline	(translation inhibitor)
Tachniplesin II	(pore former, crab)	Doxycyclin	(translation inhibitor)
Omiganan	(LL37 derivative, in clinical trials)	Streptomycin	(aminoglycoside)
Cecropin 1	(pore former, <i>Ascaris suum</i> )	Tobramicin	(aminoglycoside)
R8	(synthetic)	Cefoxitin	
Protamine	(arginine rich, herring)	Erythromcin	(macrolid)
Bactenectin 5	(proline rich, bovine)	Cloramphenicol	
PR39	(proline rich, pig)	Nalidixic acid	(gyrase inhibitor)
Pexiganan	(pore former, amphibian, in clinical trials)	Ciprofloxacin	
Indolicidin	(bovine)		