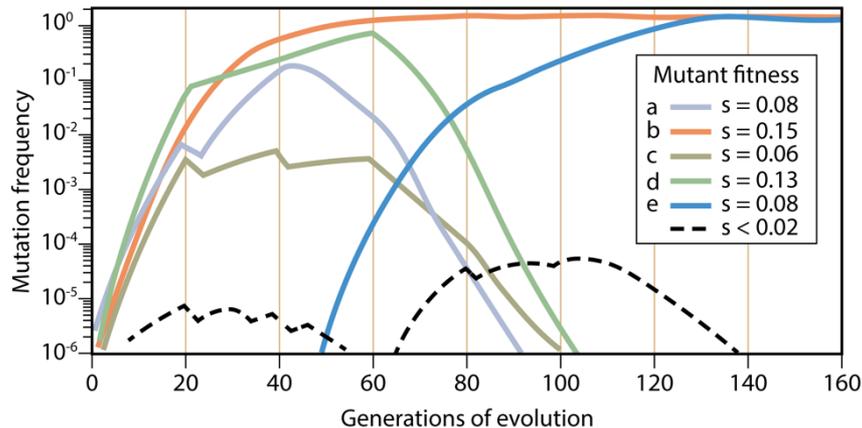


# Experimental evolution of antimicrobial resistance in *A. baumannii*

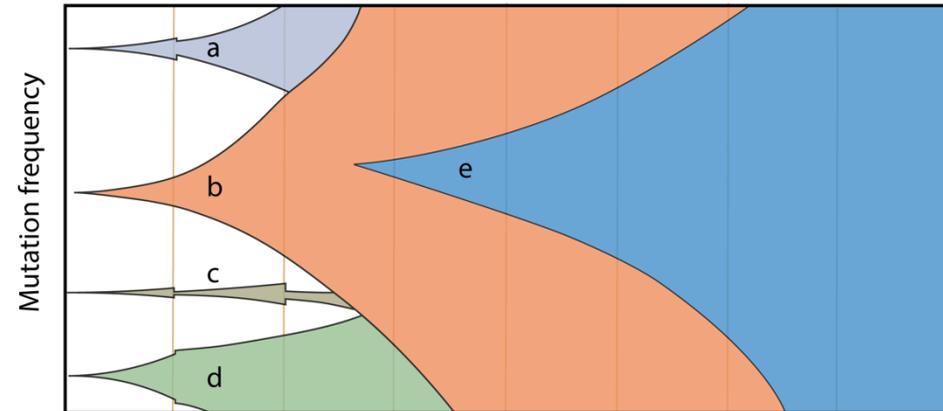
Vaughn Cooper  
Alfonso Santos-Lopez  
Michelle Scribner  
Chris Marshall

# Experimental evolution, combined with longitudinal population-wide WGS, is:

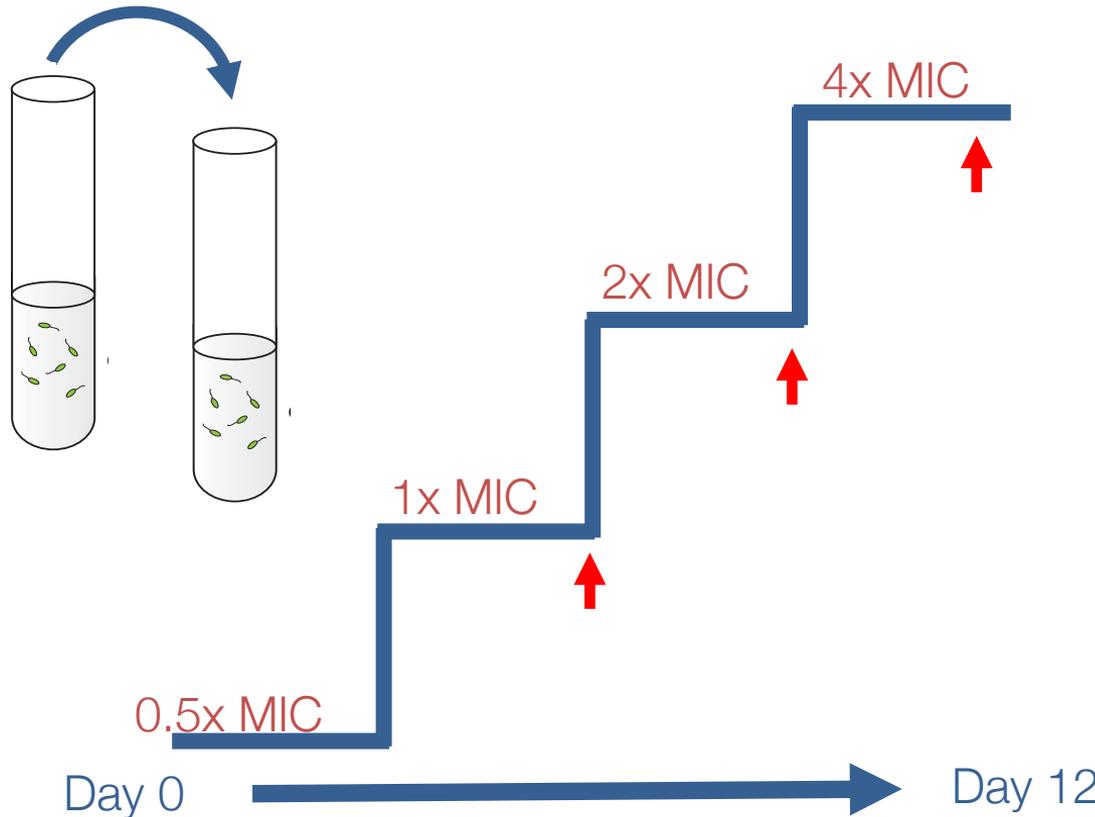
- A forward genetic screen for the **best mutations** in a given environment



- A method for quantifying relative fitness of competing genotypes



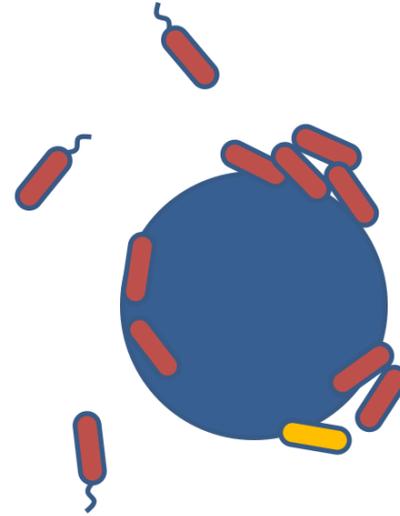
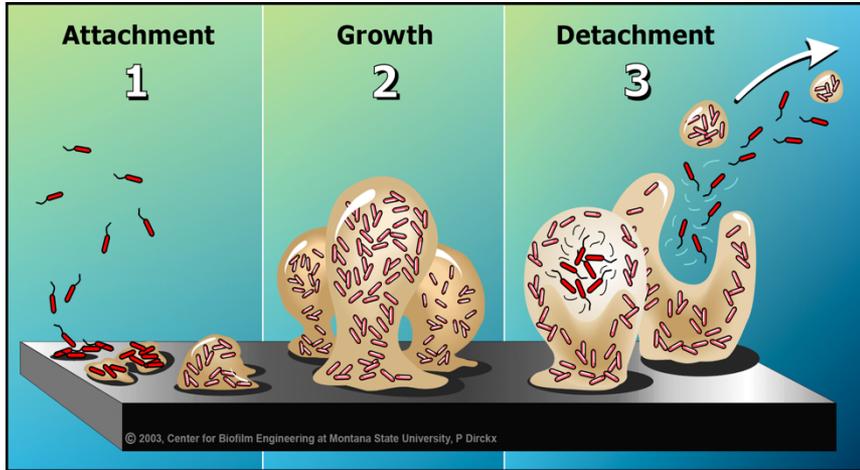
# Evolution experiments in antibiotics



MIC =  
Minimum  
Inhibitory  
Concentration

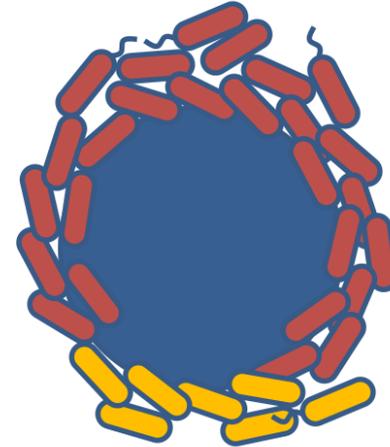
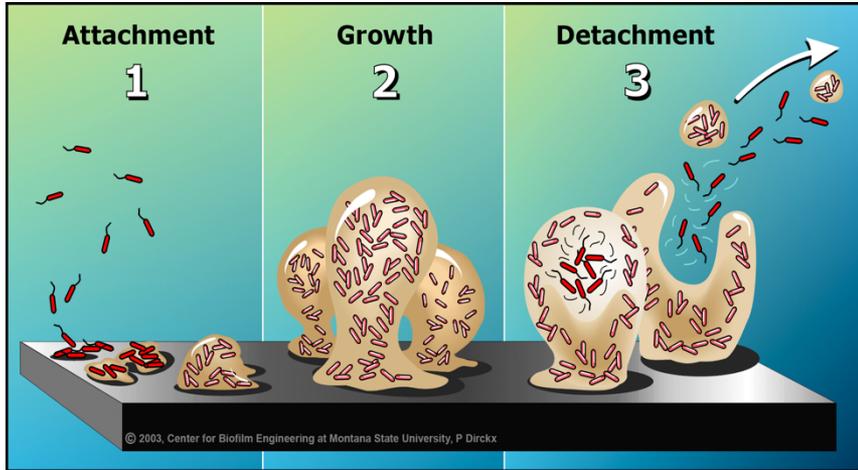
...combined  
with periodic  
whole population  
sequencing

# Bead transfer model produces strong biofilm selection



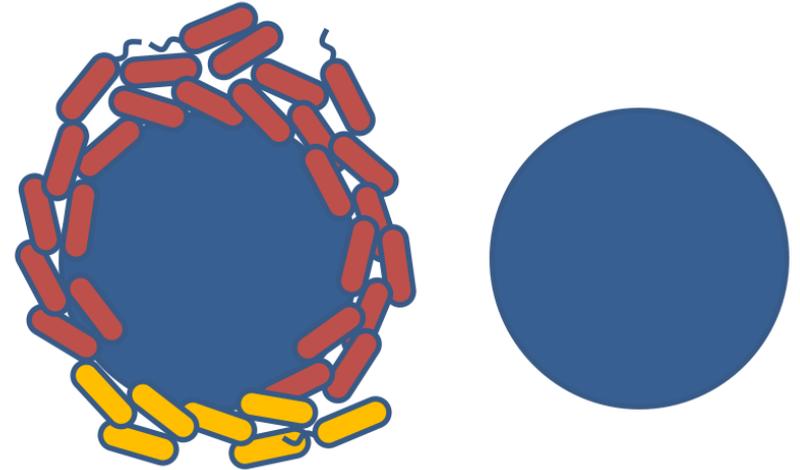
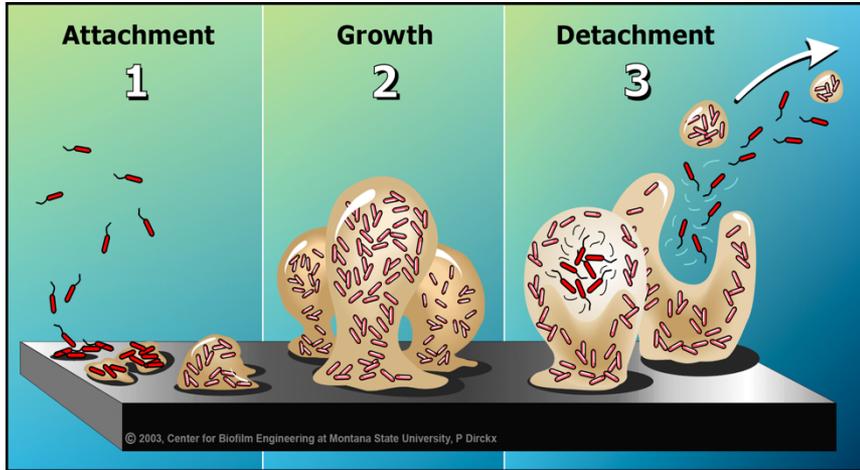
**Biofilm**

# Bead transfer model produces strong biofilm selection



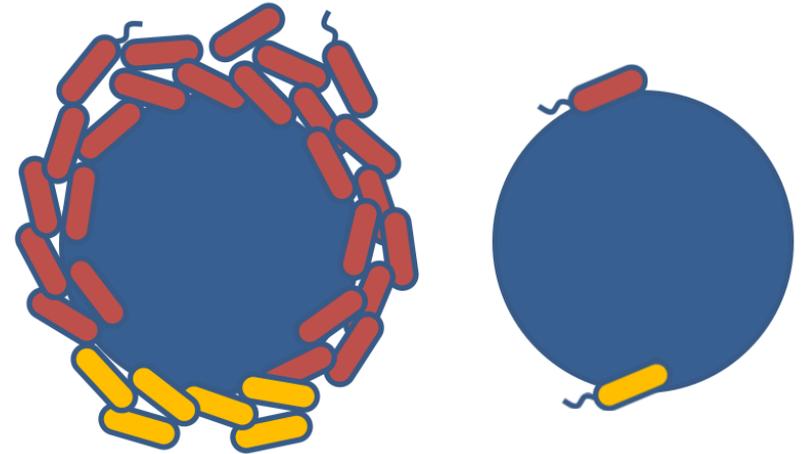
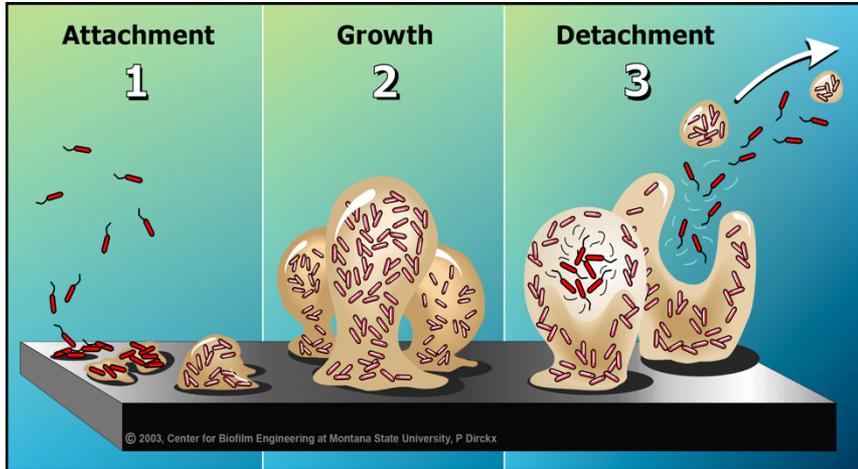
**Biofilm**

# Bead transfer model produces strong biofilm selection



**Biofilm**

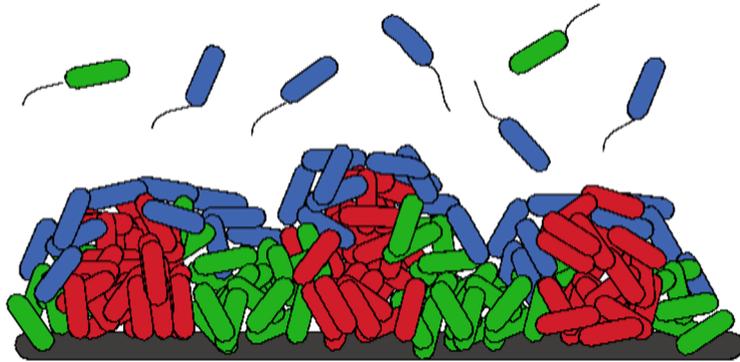
# Bead transfer model produces strong biofilm selection



**Biofilm**

See protocol description and videos at <http://evolvingstem.org>

# Biofilms can produce intrinsic tolerance to antibiotics



Matrix of exopolysaccharides, eDNA, and protein enabling cells to attach to surfaces

Hypothesis: lifestyle influences the dynamics and genetic routes to antimicrobial resistance

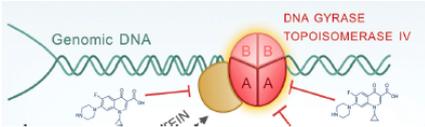


# Experimental evolution of antibiotic resistance: Design



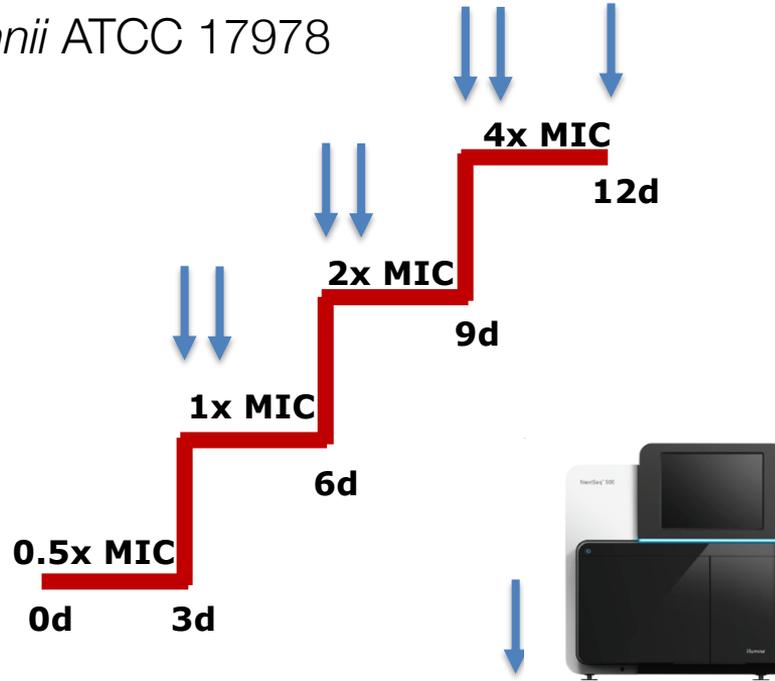
*Acinetobacter baumannii* ATCC 17978

CIP: Fluoroquinolone,  
topoisomerase inhibitor



Predictable target:  
*gyrA* S81/83L

Also: efflux-mediated





# Experimental evolution of resistance: MICs

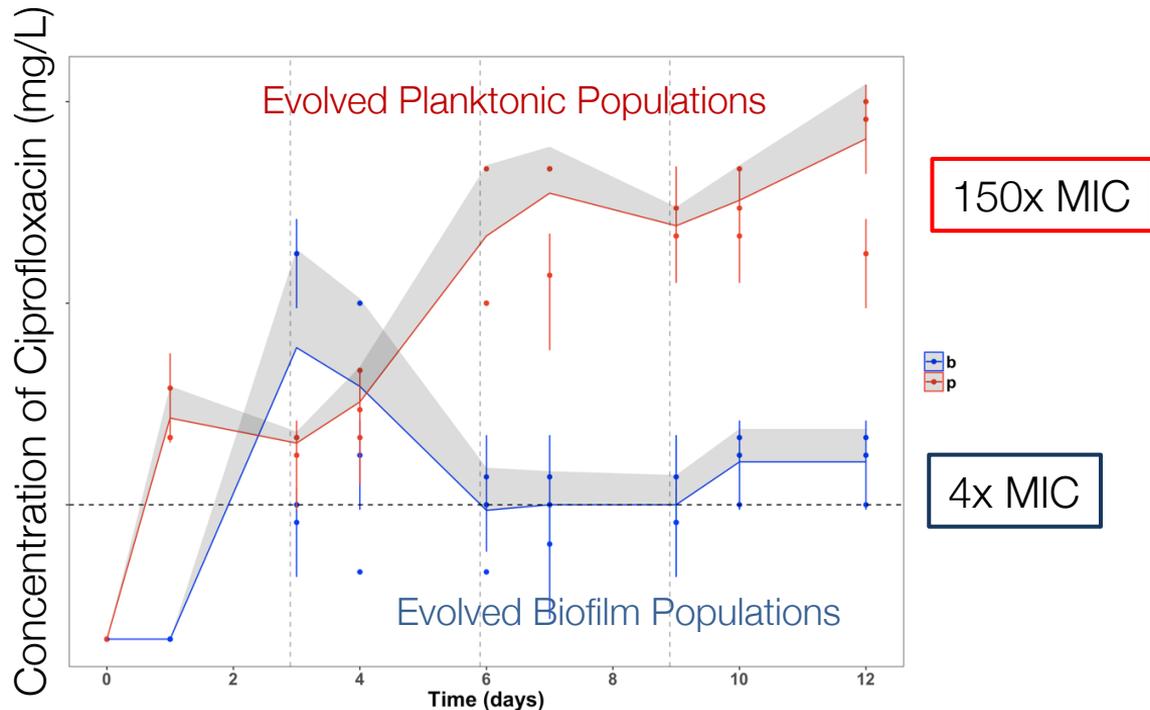
Planktonic =  
high resistance  
2+ jumps

Biofilm = one  
low resistance  
jump

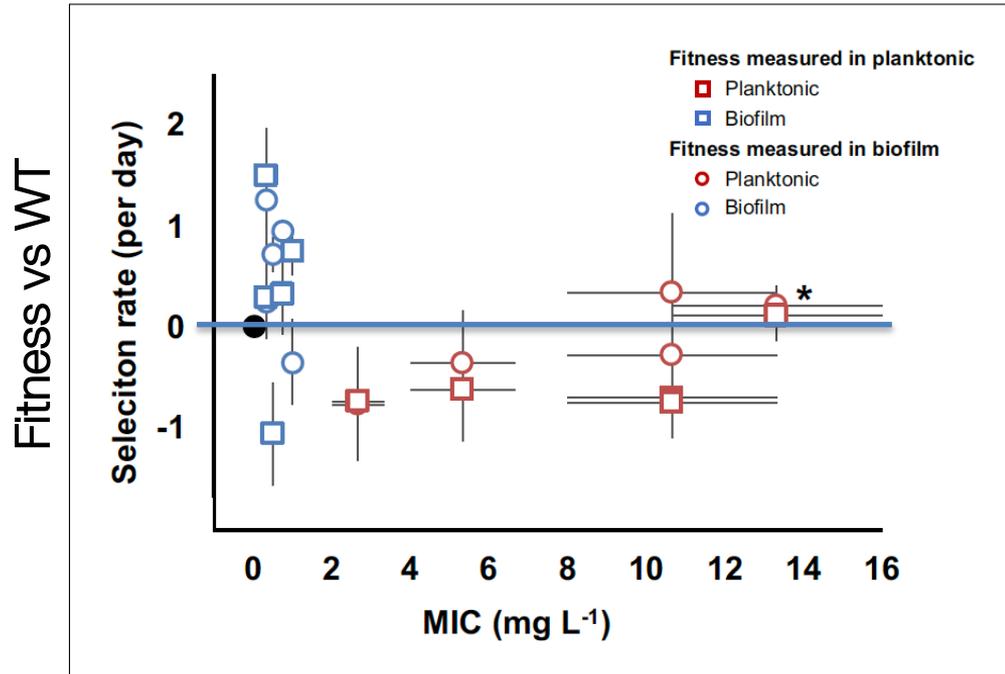
MIC increases in biofilm:

P: >1024x

B: 256x



# Evolved tradeoff between resistance and fitness





# Evolution of Antibiotic Resistance: Dynamics

**Planktonic**

Blues: *adeN*

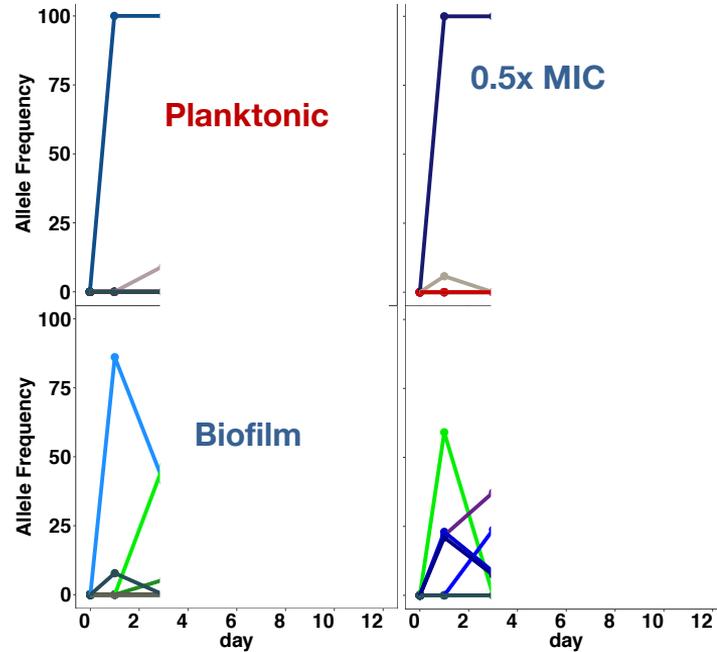
**Biofilm**

Greens: *adeL*

Blues: *adeN*

Purple: *adeS*

Ade =  
Acinetobacter  
Drug  
Efflux



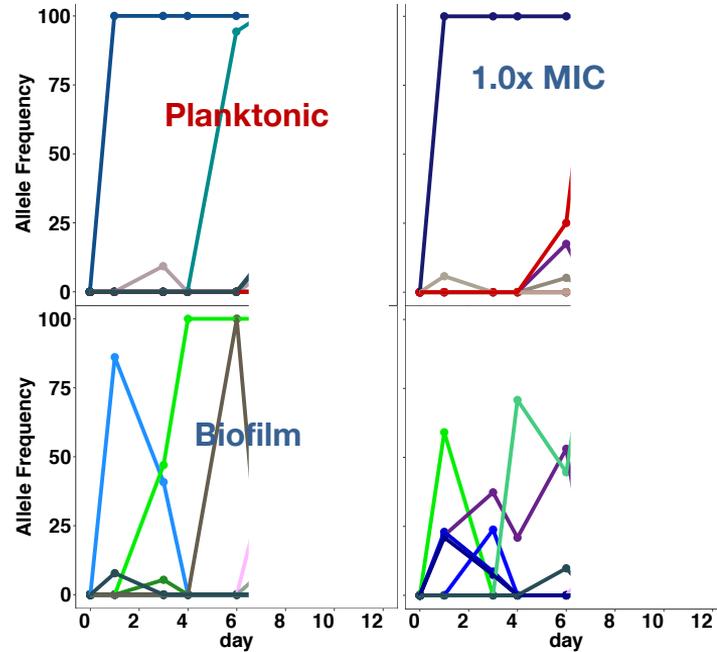


# Evolution of Antibiotic Resistance: Dynamics

**Planktonic**  
Blues: *adeN*

**Biofilm**  
Greens: *adeL*  
Blues: *adeN*  
Purple: *adeS*

Ade =  
Acinetobacter  
Drug  
Efflux





# Evolution of Antibiotic Resistance: Dynamics

**Planktonic**

Blues: *adeN*

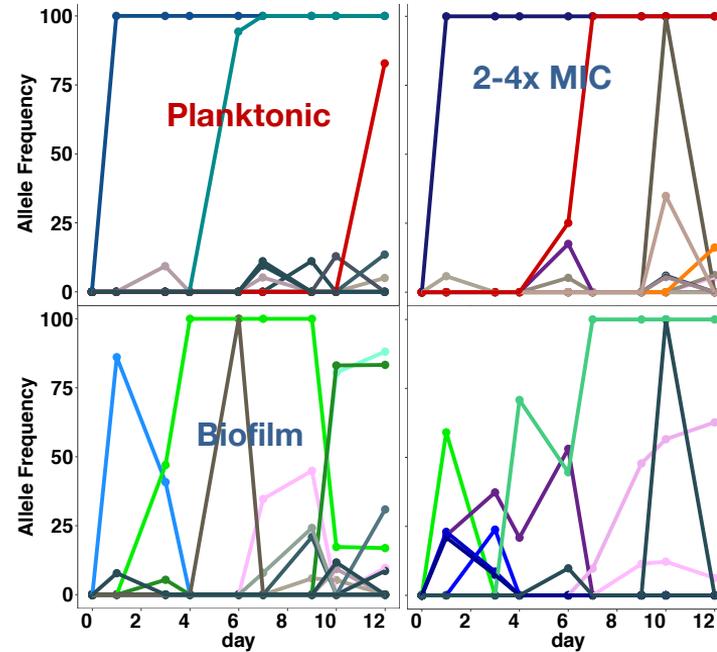
**Biofilm**

Greens: *adeL*

Blues: *adeN*

Purple: *adeS*

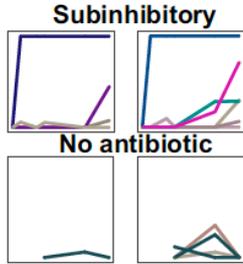
Ade =  
Acinetobacter  
Drug  
Efflux





# Evolution of Antibiotic Resistance: Dynamics

C

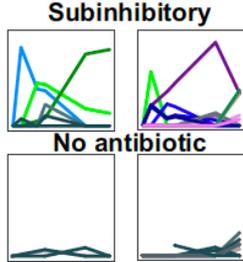


Planktonic

Blues: *adeN*

Red: *gyrA*

D



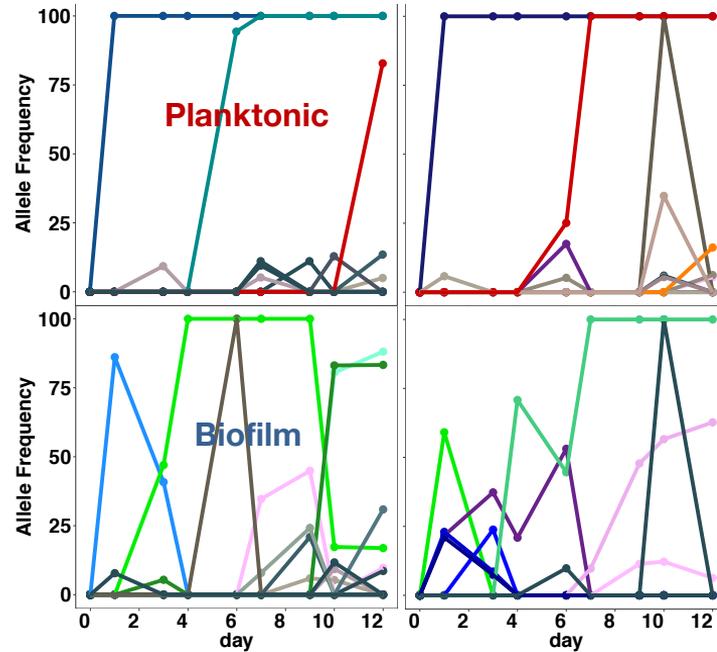
Biofilm

Greens: *adeL*

Blues: *adeN*

Purple: *adeS*

Ade =  
Acinetobacter  
Drug  
Efflux

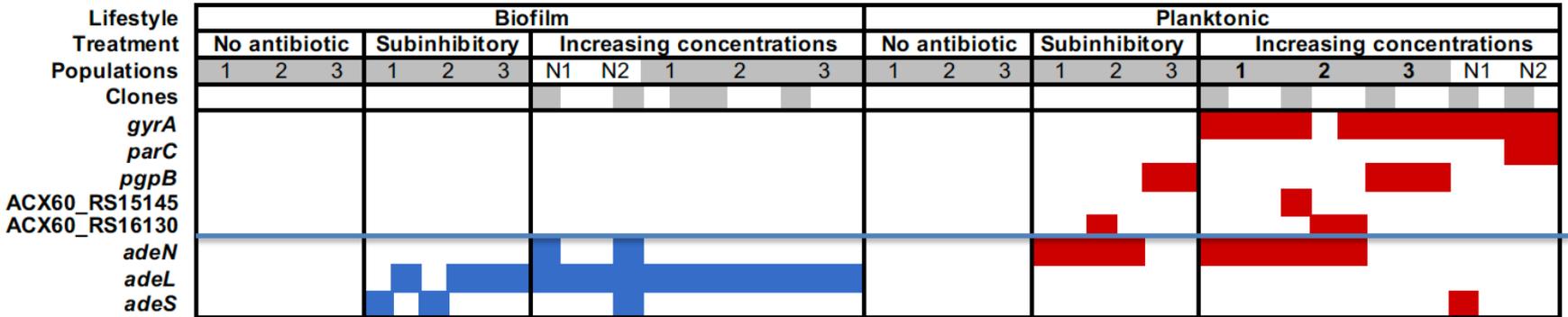


More  
diversity

More  
diversity



# Evolution of Antibiotic Resistance: Genotypes differ w/ Lifestyle

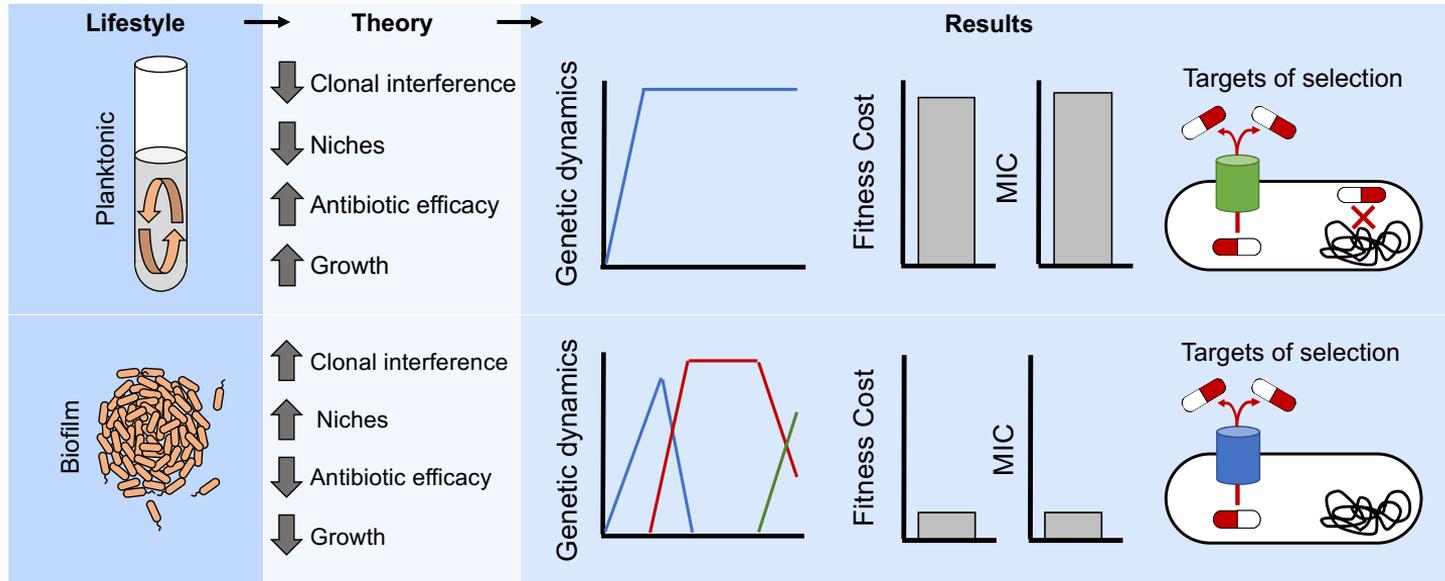


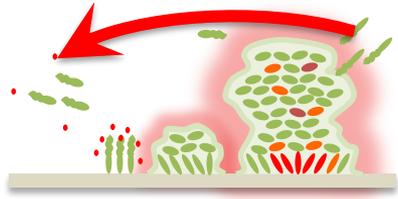
*adeL* and *adeS* dominate

*gyrA* and *adeN* dominate



# Selection (predictably) acts on different mutations with different effects in different environments





## Why resistance evolution in biofilms is different

1. Environmental structure increases genetic diversity
2. Greater ecological heterogeneity = multiple niches
3. Biofilms provide intrinsic protection to external stresses like antibiotics

### **Parallel evolution of tobramycin resistance across species and environments**

Michelle R. Scribner,  Alfonso Santos-Lopez,  Christopher W. Marshall, Christopher Deitrick,  Vaughn S. Cooper

**doi:** <https://doi.org/10.1101/758979>

Now accepted for publication in *mBio*

Related project about aminoglycoside resistance evolution in *A. baumannii* and *P. aeruginosa*, see: