Chemical Biology approach to identification of a small-molecule inhibitor of BamA function



March 2020 D. M. Rothman on behalf of team

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Gram-negative Challenge

The Threat of Antibiotic Resistance in the United States

Antibiotic resistance—when germs (bacteria, fungi) develop the ability to defeat the antibiotics designed to kill them—is one of the greatest global health challenges of modern time.

New National Estimate*

Each year, antibiotic-resistant bacteria and fungi cause at least an estimated:







223,900

Clostridioides difficile is

antibiotic resistance:

related to antibiotic use and

New Antibiotic Resistance Threats List

Updated urgent, serious, and concerning threats-totaling 18

5 urgent threats

2 new threats

Watch List with 3 threats



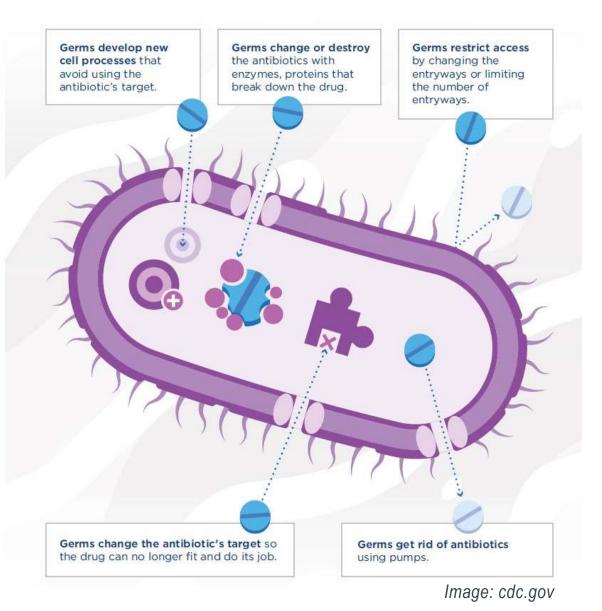
Antibiotic resistance remains a significant One Health problem, affecting humans, animals, and the environment. Data show infection prevention and control is saving lives—especially in hospitals—but threats may undermine this progress without continued aggressive action now. Antibiotic resistance remains an increasing threat to public health and three of the five urgent threat pathogens are gram-negative bacteria



Images: cdc.gov

INVENTING FOR LIFE

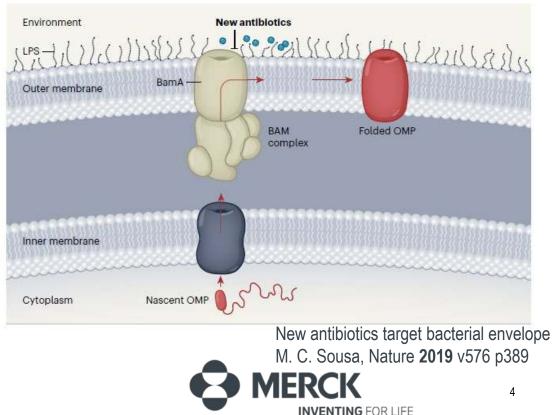
Gram-negative Challenge



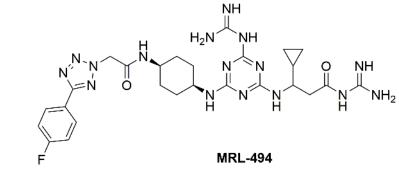
Gram-negative antibiotic discovery is additionally challenged by the Outer Membrane (OM)

- Asymmetric bilayer (Outer Leaflet = lipopolysaccharides, Inner Leaflet = phospholipids)
- MDR efflux pumps

Targeting the OM as a strategy

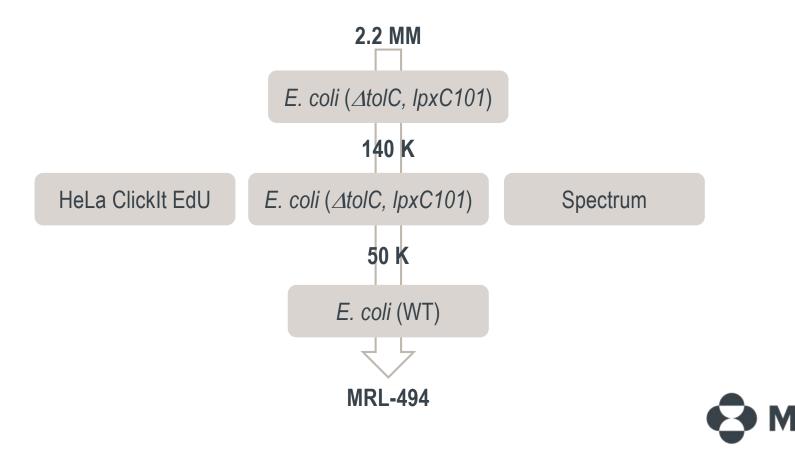


Approach and Identification of MRL-494

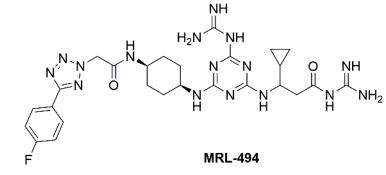


NVENTING FOR LIFE

Hypothesis: compounds with similar activity on efflux and permeability competent and impaired cells may evade efflux and the OM permeability barrier due to chemical nature and/or the surface location of their target



Profiling MRL-494



Spectrum activity of MRL-494

Classification	Description	MIC (µM)
Gram-negative	E. coli (WT)	25
	E. coli (∆tolC)	25
	E. coli (∆tolC, lpxC101)	25
	K. pneumoniae	100
	A. baumannii (WT)	200
	A. baumannii (∆lpxC)	200
	P. aeruginosa (WT)	100
	P. aeruginosa (efflux deficient)	100
Gram-positive	Staphylococcus aureus (methicillin-resistant)	12.5
	Bacillus subtilis rpoB18	25
HeLa	Mammalian – ClickIt EdU (EC50)	> 99

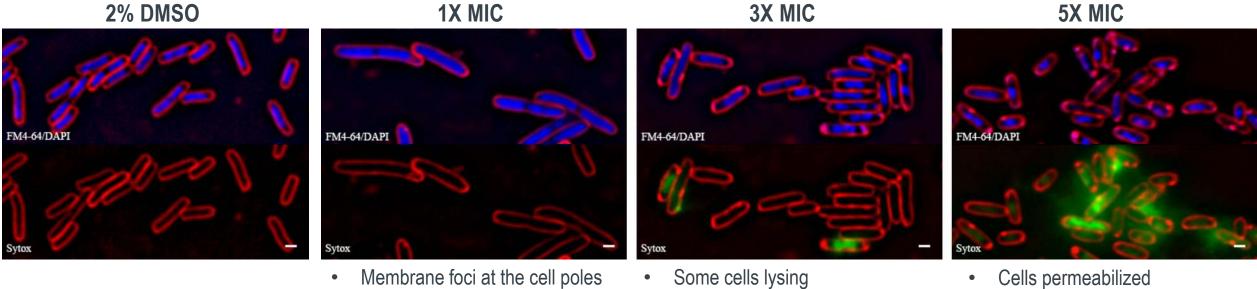
Not efflux substrate No permeability issue



Initiating MOA Deconvolution

FM4-64: membrane **DAPI:** nucleic acid Sytox: nucleic acid, non-permeable 2hr compound exposure

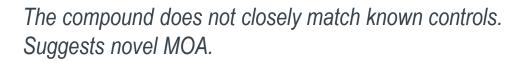
Bacterial Cytological Profiling (Linnaeus)



Some abnormally long

Some cells lysing

- Cells permeabilized
- Cells lysing •





MRL-494 impacts OM and decreases OMP abundance

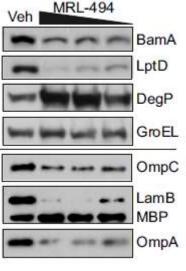
MRL-494 potentiates rifampicin

Condition	<i>E. coli</i> MIC (uM)		
Condition	WT	lpxC101	
Rifampicin	25.000	< 0.049	
Rifampicin + 6.25 uM MRL-494	0.195		

Rifampicin: RNA polymerase inhibitor which does not penetrate OM

IpxC101: mutation which decreases LPS on OM and potentiates rifampicin activity

MRL-494 decreases OMPs



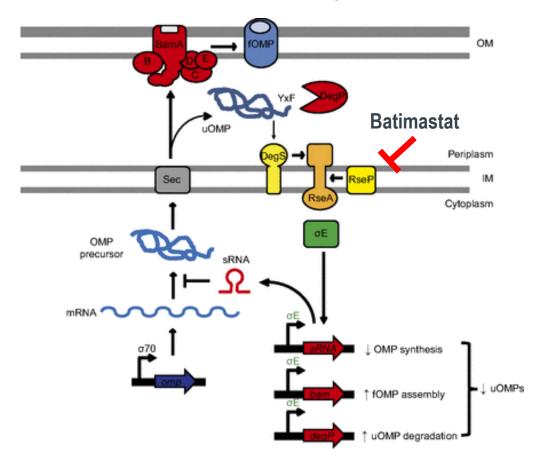
1.5h treatment

Observe increase in DegP Indication of extracellular envelope stress



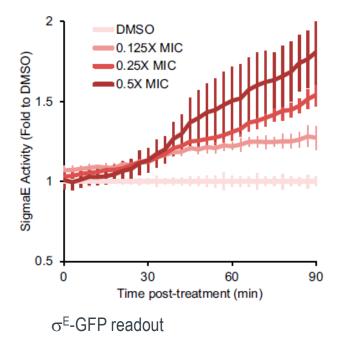
MRL-494 increases stress response

Batimastat targets RseP and inhibits the σE pathway



PNAS 2018 115 (28) E6614

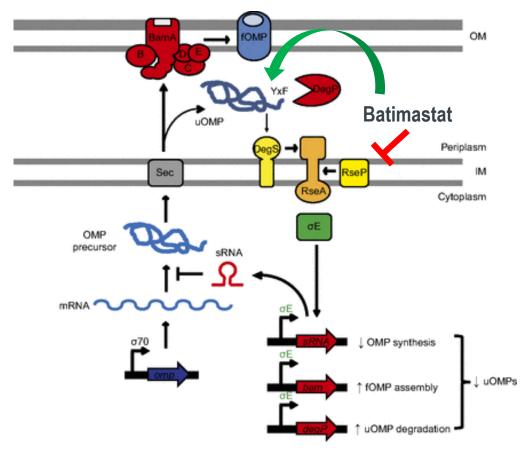
MRL-494 increases σ^{E} activity





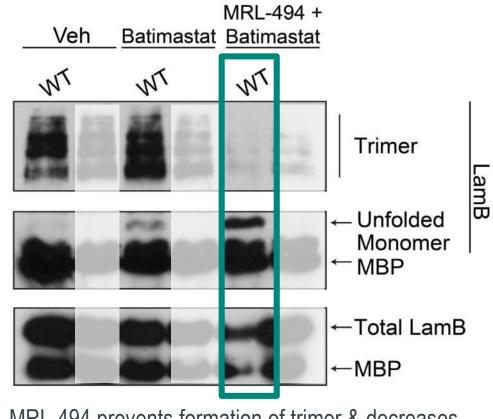
MRL-494 inhibits OMP biogenesis

Batimastat targets RseP and inhibits the σE pathway



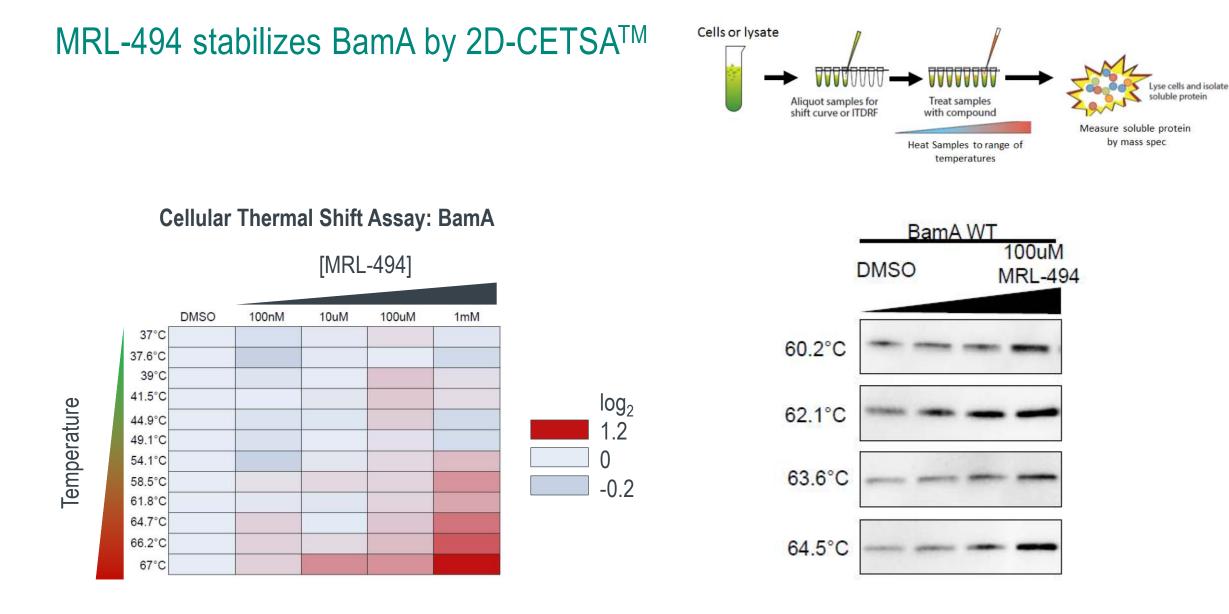
PNAS 2018 115 (28) E6614

Batimastat potentiates MRL-494



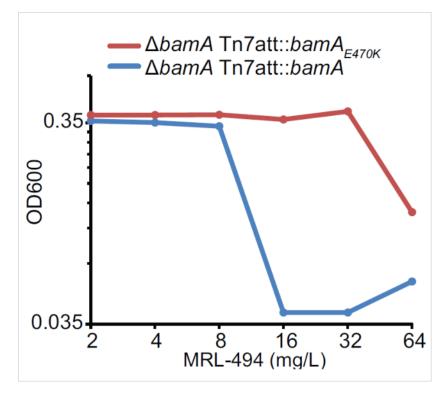
- MRL-494 prevents formation of trimer & decreases total amount of LamB
- Batimastat increases accumulation of unfolded OMP



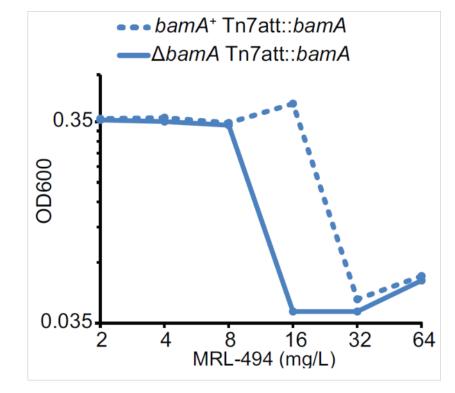


BamA genetic status impacts resistance to MRL-494

BamA^{E470K} confers resistance

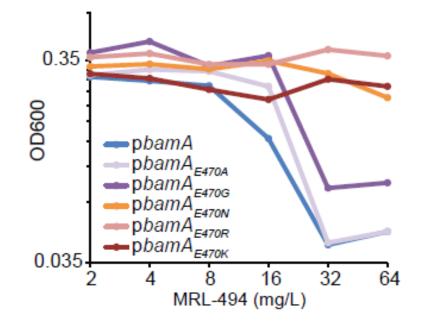


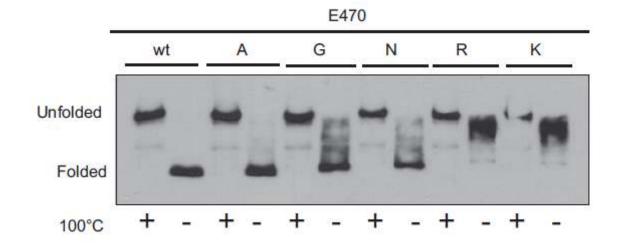
BamA WT OE confers resistance





Observed 470 amino acid charge impacts MRL-494 activity and BamA stability

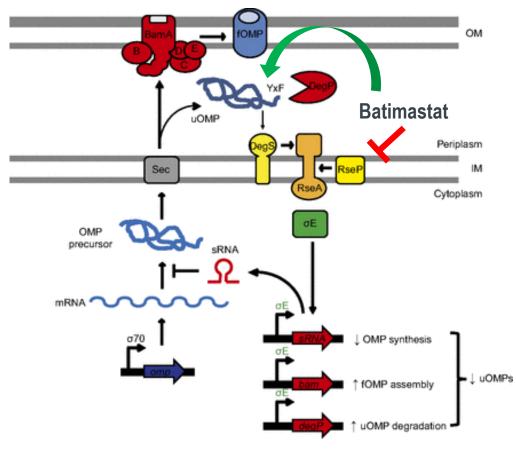






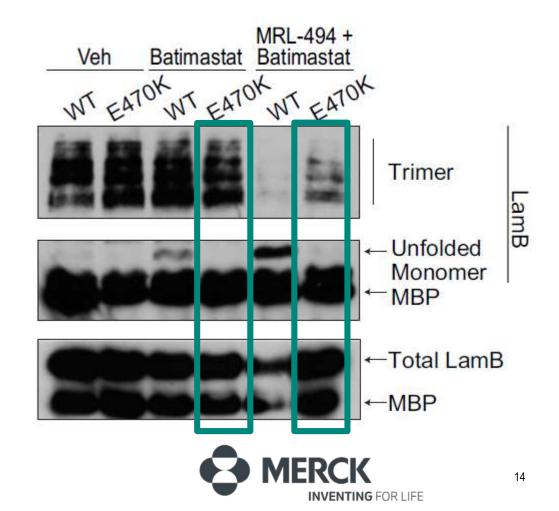


Batimastat targets RseP and inhibits the σE pathway

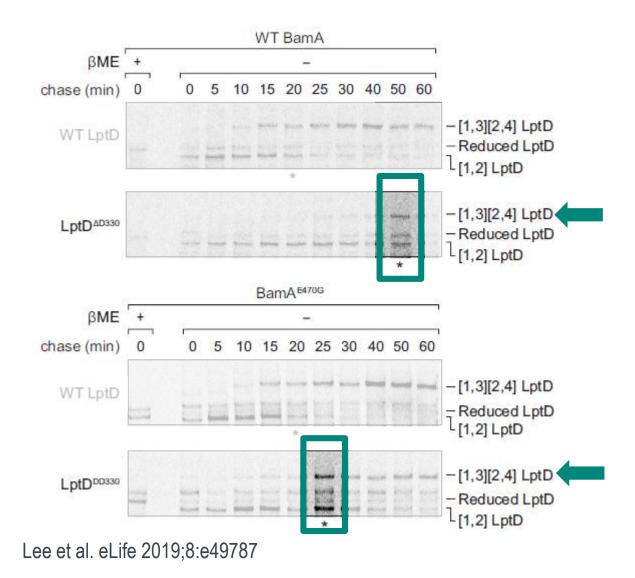


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E470K mutant suppresses batimastat/MRL-494



BamA^{E470} mutation may act as hypermorph



Pulse chase experiment recently reported from Kahne, et. al.

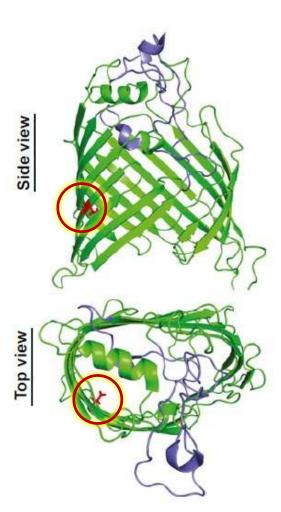
- * = approximately 50% of the LptD has converted to the mature form, containing the [1,3][2,4] disulfide bond configuration
- BamA WT ~ 50 min BamA^{E470G} ~ 25 min



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- ✓ MRL-494 identified as tool which functionally inhibits BamA in Gram-negative bacteria
- ✓ BamA^{E470K} identified as mutant which demonstrates MIC shift for MRL-494
- BamA^{E470K} does not have impaired function in the assembly of wild-type OMPs under laboratory conditions
- ✓ BamA^{E470K} does have both altered conformation and activity
- Correlates with resistance to MRL-494 Neither the conformational change nor the altered activity sufficient to explain the MRL-494 resistance
- Recent publication demonstrates opportunity for how MRL-494 could be a mechanistic tool to inform on new biology – how would different mutations impact MRL-494 activity?





THANK YOU

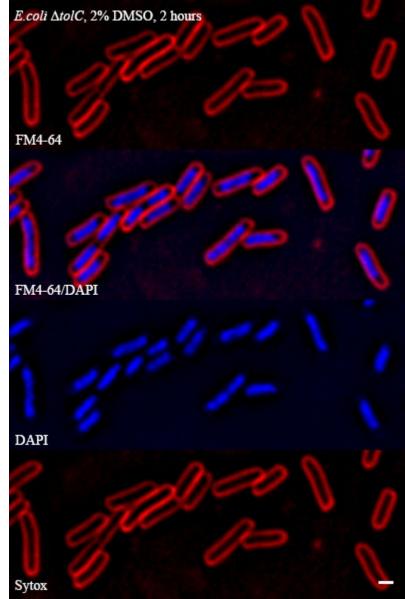


Bacterial Cytological Profiling Linnaeus

- For BCP, we grew samples of *E.coli* ∆tolC in rich media (LB) at 30°C to an OD₆₀₀ ~0.15. These cultures were then split and each sample treated with the appropriate concentration of test compound.
- We collected samples after 30 minutes or 2 hours of exposure, stained and then imaged them.
- Below, we have included images showing the predominant phenotype produced by each of the test compounds at 2 hours of exposure.

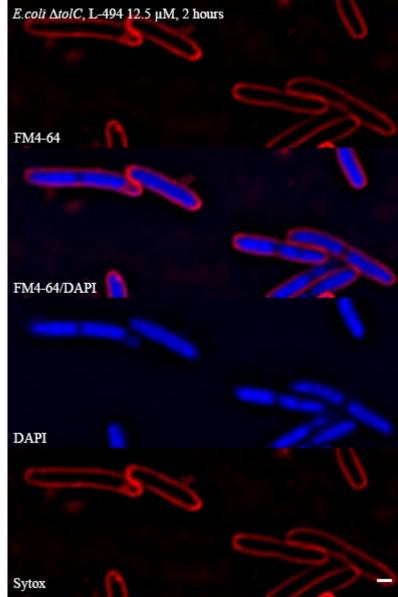
	Merck Δ <i>toIC</i> MIC (μM)	Linnaeus ∆ <i>tolC</i> MIC (µM)
L-494	12.5	12.5

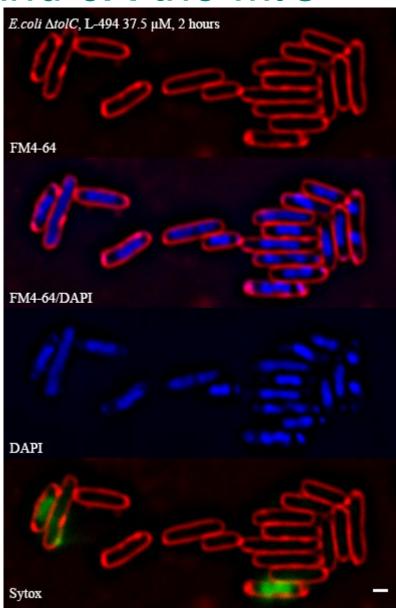
2% DMSO Control

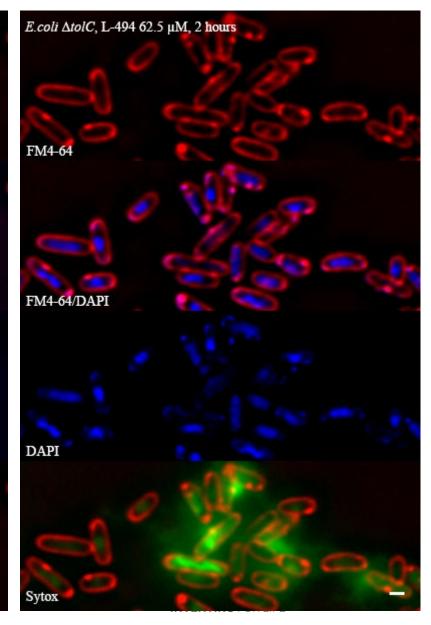




L-494 at 1X, 3X and 5X the MIC

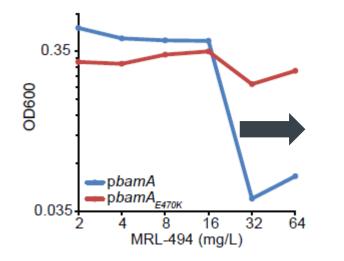






BamA^{E470K} shifts MRL-494 potency

BamA^{E470K} shifts MIC to solubility limit



BamA^{E470K} shifts MIC in ∆bamB

