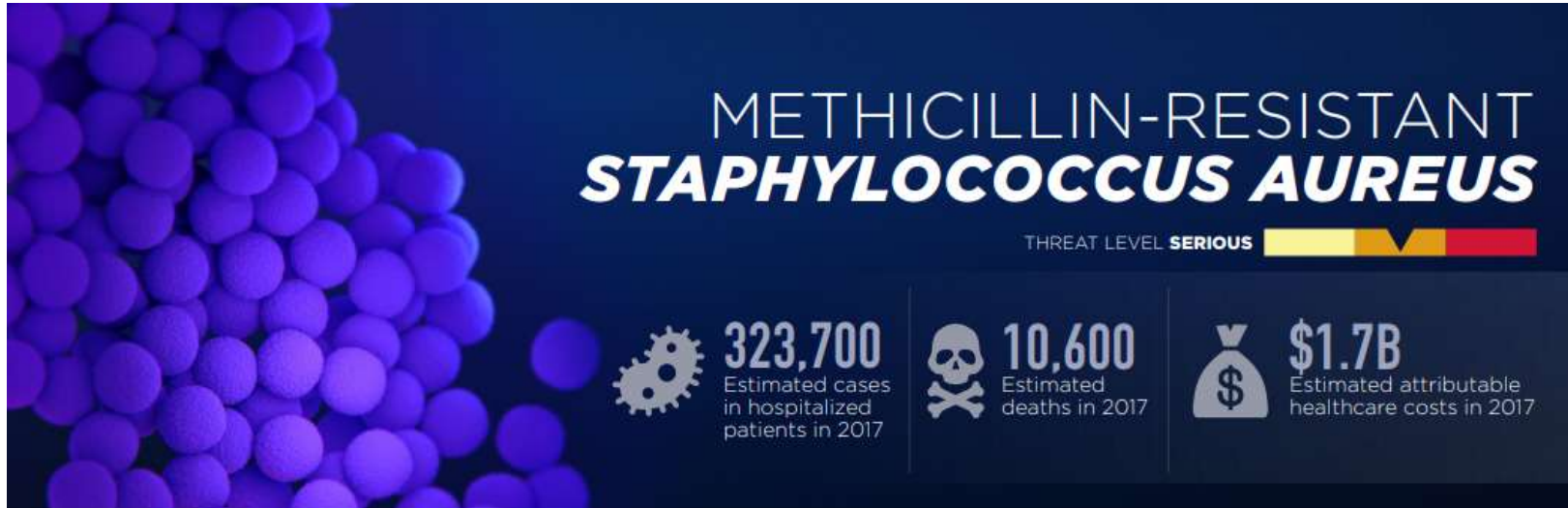


Mutations that inactivate the Tricarboxylic Acid Cycle in
Staphylococcus aureus arise during persistent MRSA bacteremia

Edwin Chen MD PhD
Post-doctoral Research Fellow
Culyba Lab
Division of Infectious Diseases
Department of Medicine
University of Pittsburgh

Invasive MRSA infections continues to be a present threat



Persistent MRSA bacteremia is common with high morbidity and mortality despite appropriate antibiotics.

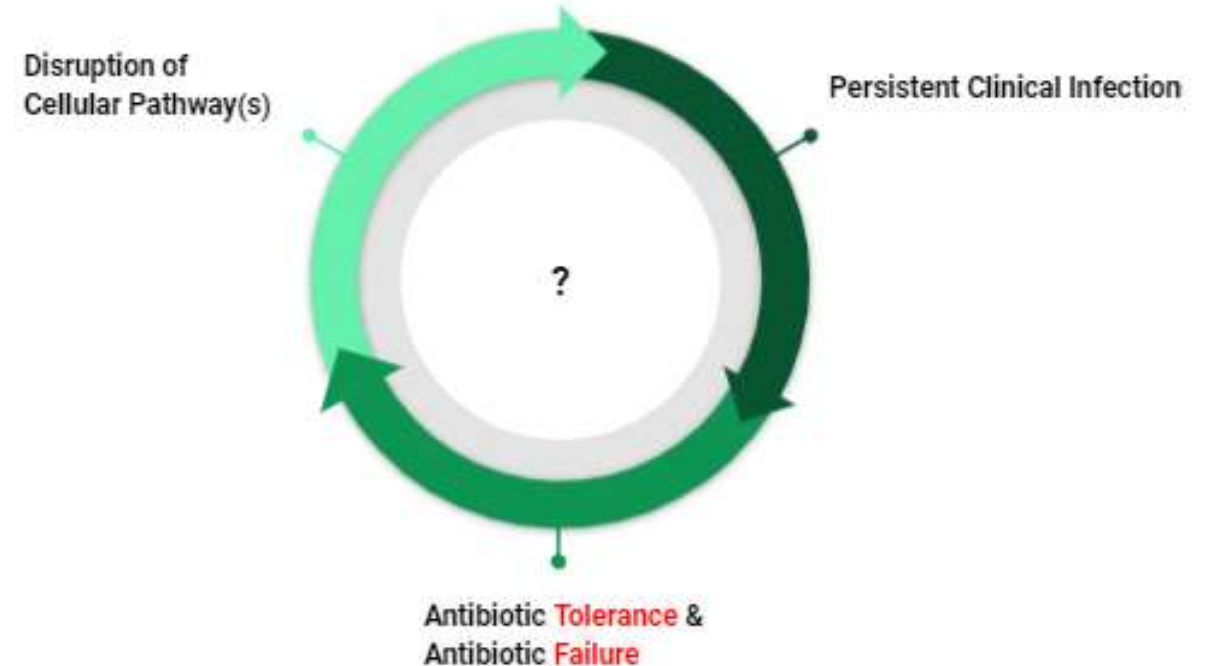
There is great interest in understanding the mechanism(s) behind persistent MRSA infections.

Persistent MRSA infections are associated with perturbations in cellular pathways and antibiotic tolerance

Clinically persistent *S. aureus* infections have been associated with alterations in cellular pathways and **antibiotic tolerance**.

These persistent infections have been linked to phenotypes associated with **antibiotic failure** despite appropriate antibiotic choice.

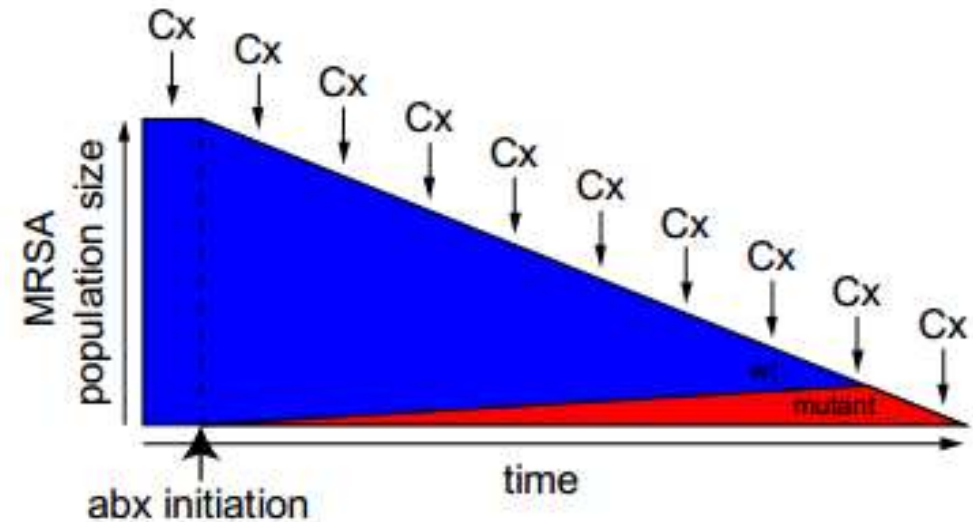
- Small Colony Variants (SCVs)
- Antibiotic “persister” sub-populations



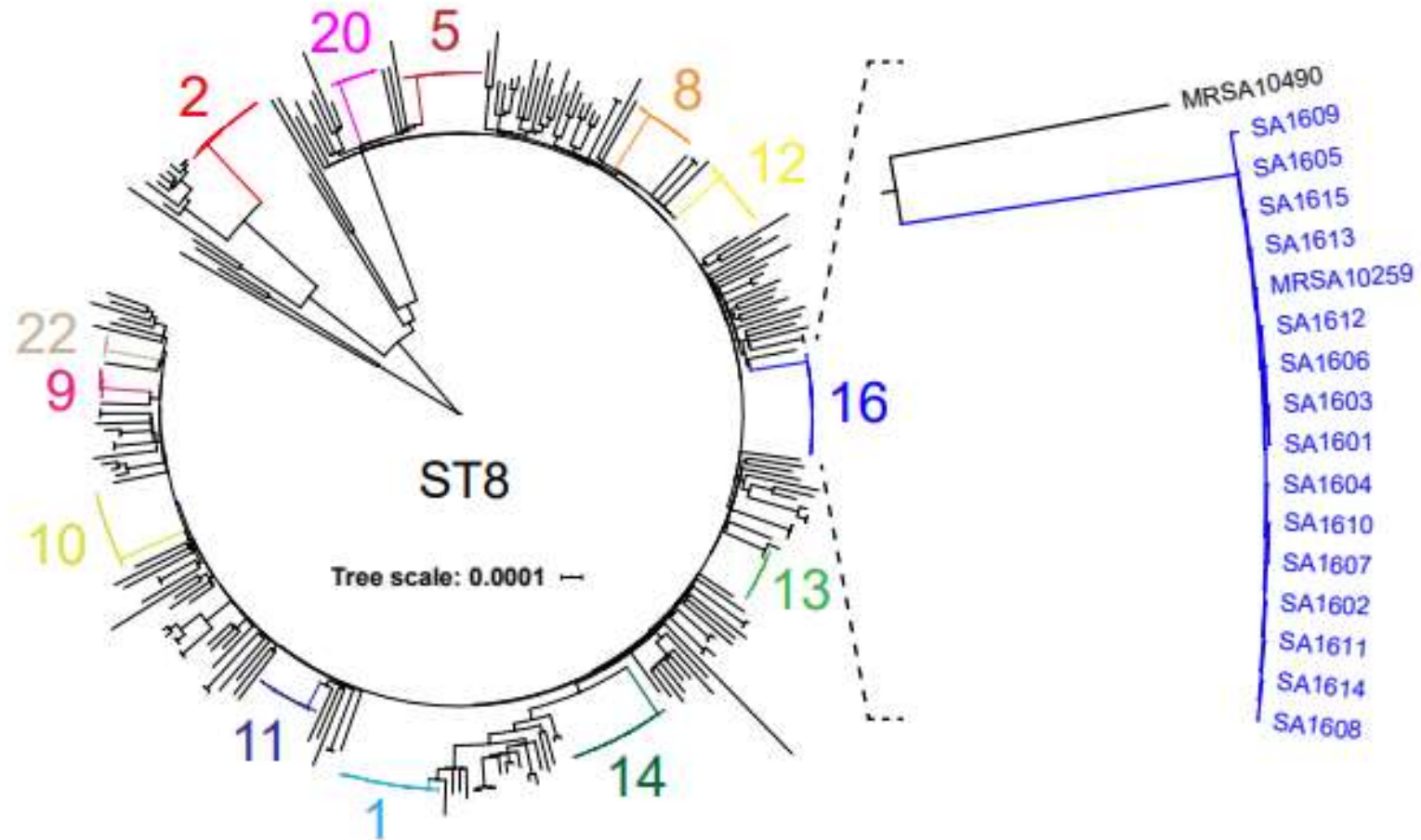
A forward genetic screen to search for within-host evolution of antibiotic tolerance

Sequenced whole genomes of 206 serially positive MRSA blood cultures from 20 patients with persistent clinical infections.

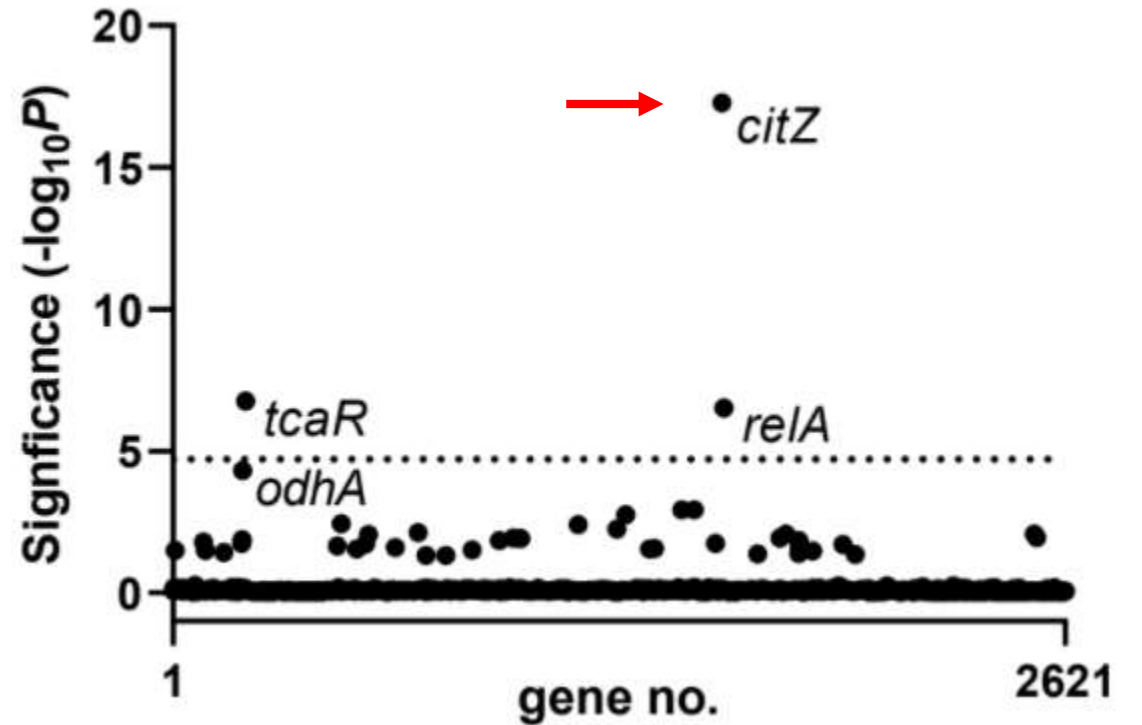
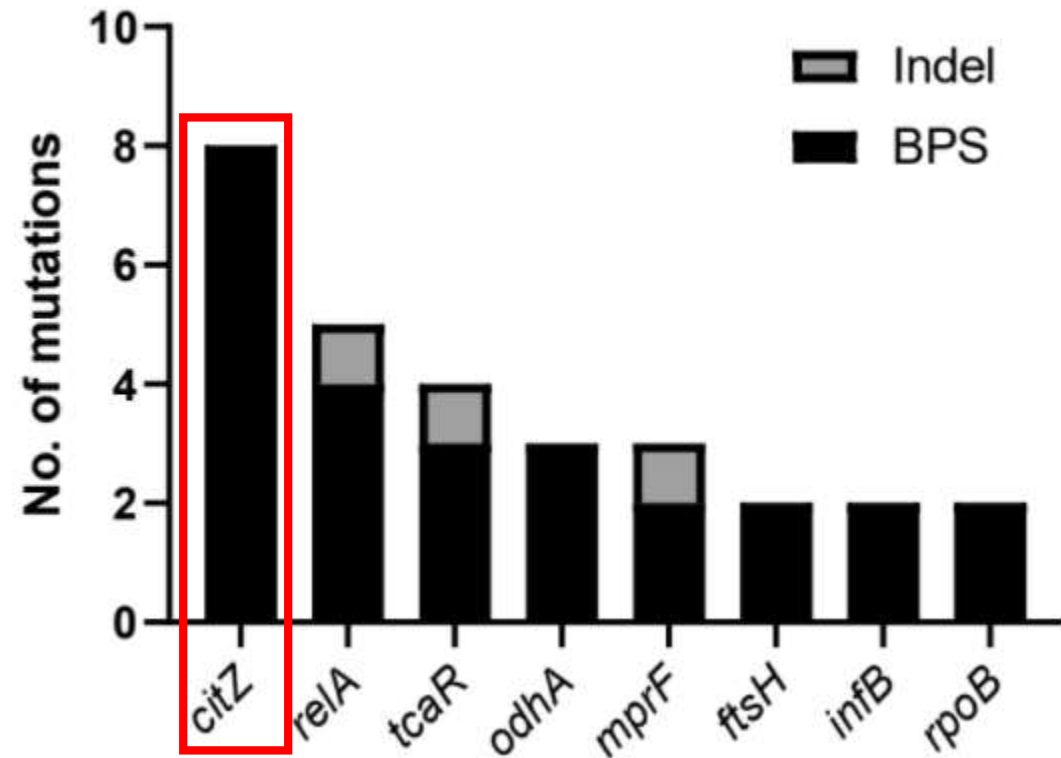
Hypothesis that continuous antibiotic exposure will give rise to mutations that convey antibiotic tolerance through within-host evolution.



MRSA from different patients are genetically distinct



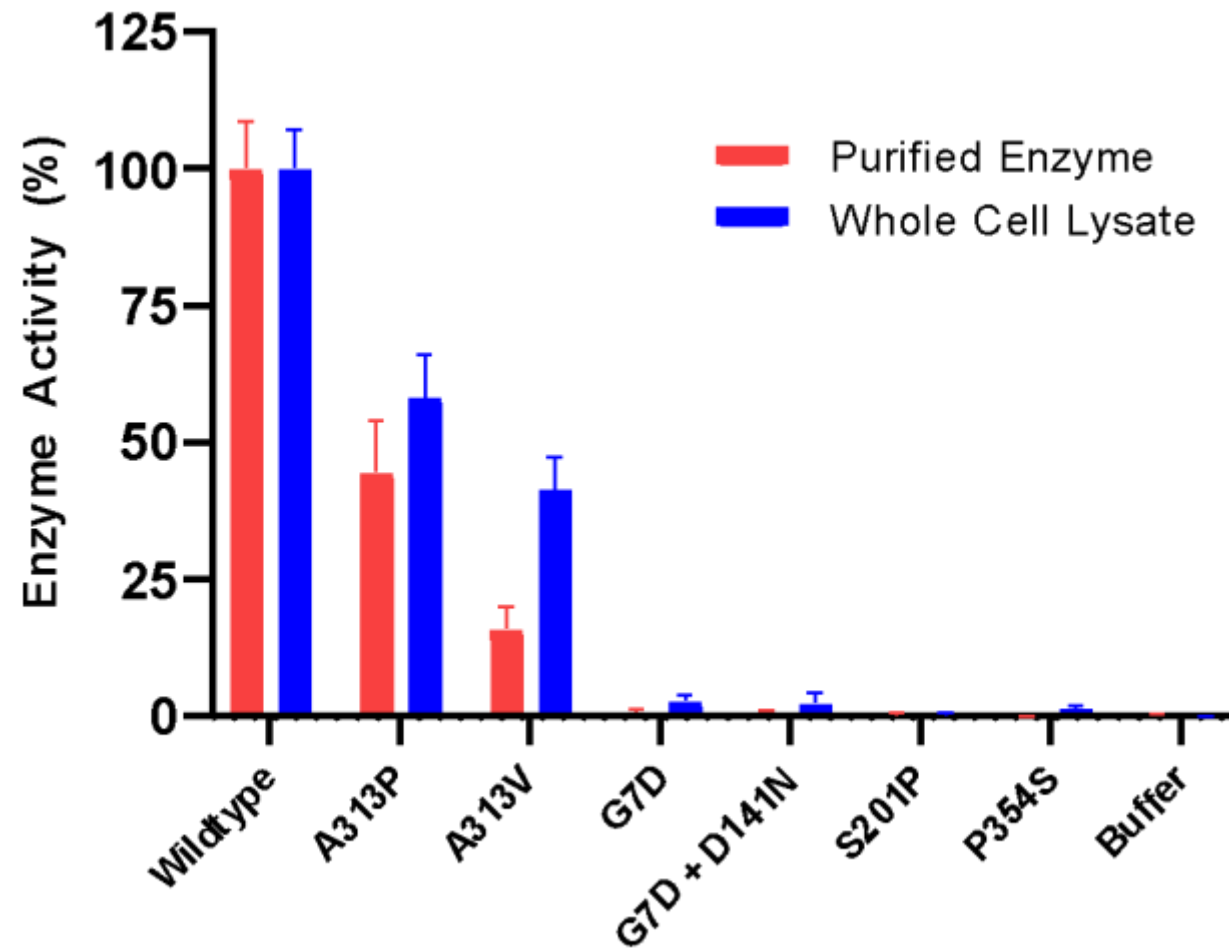
Convergent evolution of TCA cycle gene *citZ*



Key Questions

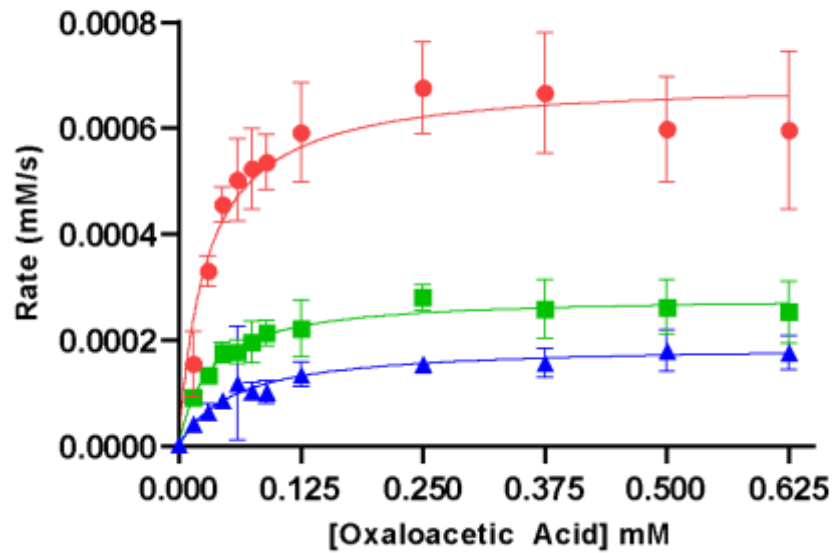
1. What are the effects of the *citZ* mutations on enzyme function?
2. What role do these mutations play in clinically persistent MRSA infections?

citZ mutations decrease citrate synthase enzyme activity

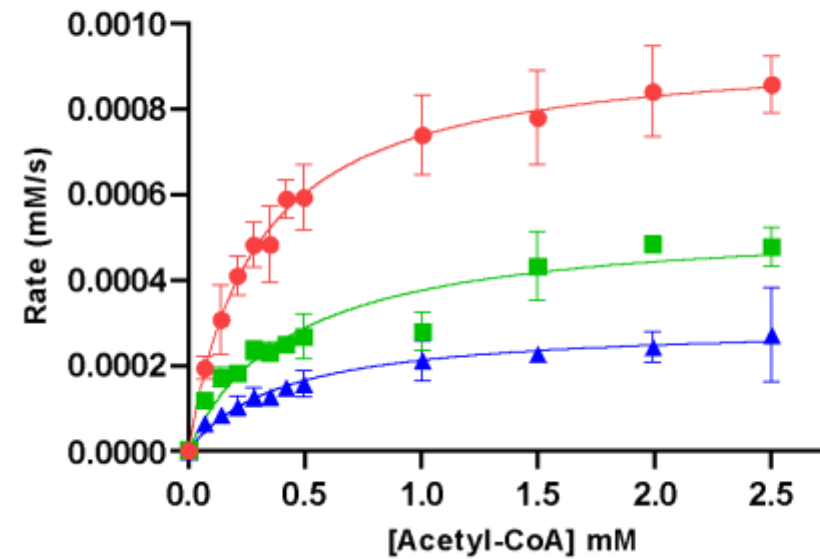


citZ mutations decrease citrate synthase enzyme activity

Acetyl-CoA + Oxaloacetic Acid \rightarrow Citrate + CoA-SH
Fixed Acetyl-CoA (0.3 mM), Variable Oxaloacetic Acid (0-0.625 mM)

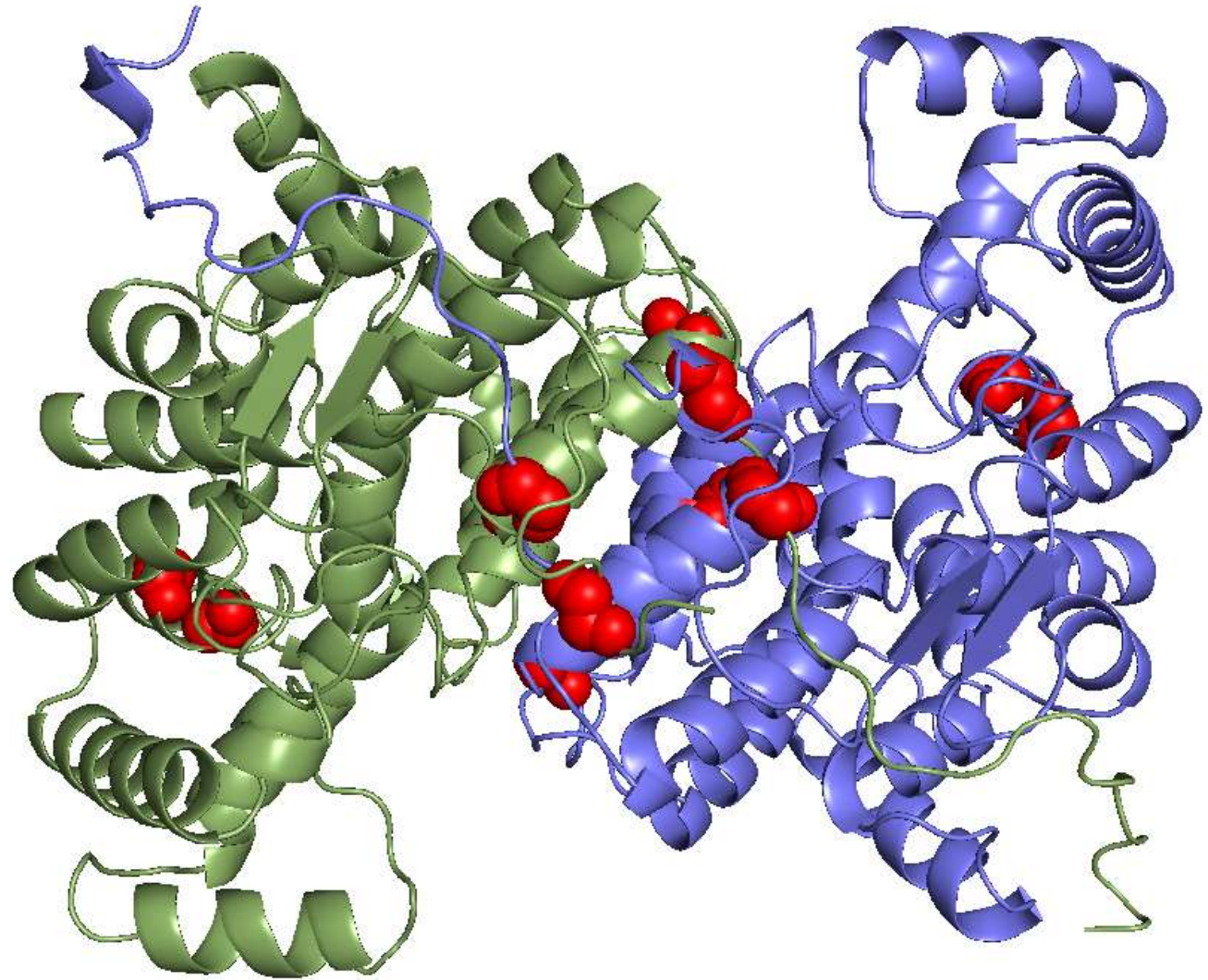


Acetyl-CoA + Oxaloacetic Acid \rightarrow Citrate + CoA-SH
Variable Acetyl-CoA (0-2.5 mM), Fixed Oxaloacetic Acid (0.5 mM)

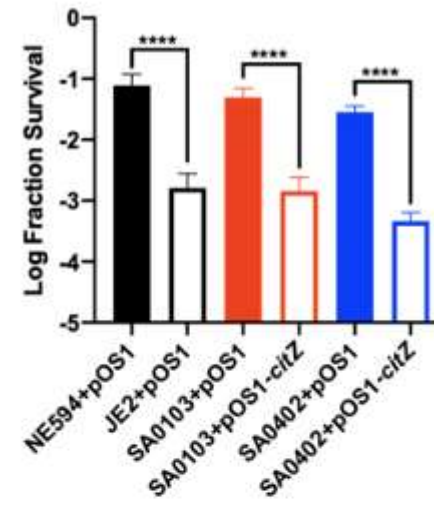
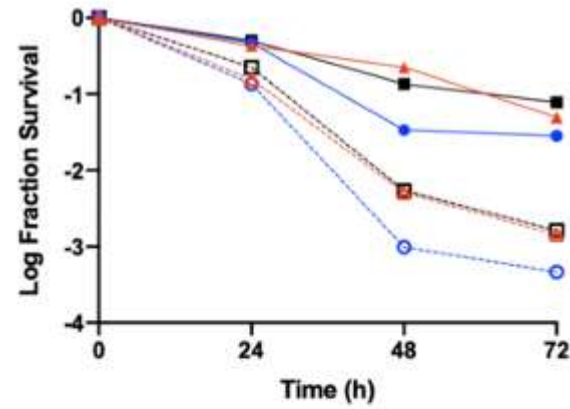


● WT ■ A313P ▲ A313V

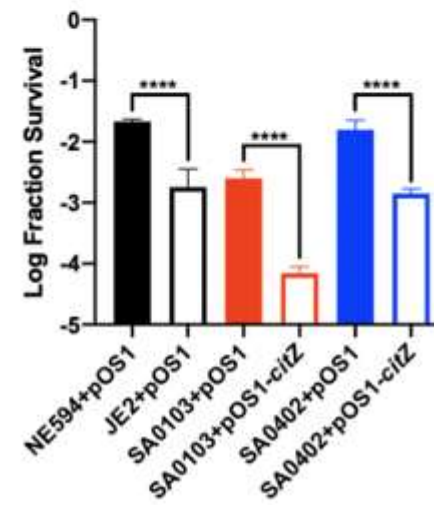
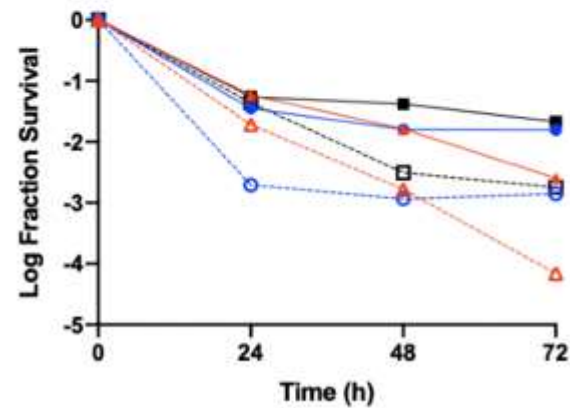
Citrate synthase mutations disrupt functional dimerization and destabilize α -helix packing



Clinical strains with *citZ* mutations have increased antibiotic tolerance



Ceftaroline



Daptomycin

Conclusions

- Convergent evolution of TCA cycle (*citZ*, *odhA*) and Stringent Response (*rel*) genes occur during persistent MRSA bacteremia.
- The strongest signal occurred in *citZ*, where the mutations caused loss of citrate synthase activity.
- Clinical strains harboring *citZ* mutations demonstrate increased antibiotic tolerance.

∴ Convergent evolution of antibiotic tolerance occurs frequently in MRSA bacteremia.

Acknowledgements

Culyba Lab

Matthew Culyba
Marla Shaffer
Mitra Eghbal

Richardson Lab

Aimee Stephens

Doi Lab

Christi McElheny
Erin Fowler

Shields Lab

Ryan Shields

Sluis-Cremer Lab

John Barnard

Harrison Lab

Mustapha Mustapha
Marissa Griffith
Vatsala Srinivasa

Urish Lab

Nguyen Lab

Van Tyne Lab

Dimitrov Lab

Division of Infectious Disease

Department of Medicine

University of Pittsburgh Medical Center

