



Investigating the role of fused *msrA/B* and *clpX* in the resistance to cell-wall antibiotics in *Bacillus anthracis* Sterne



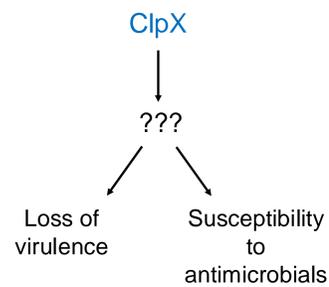
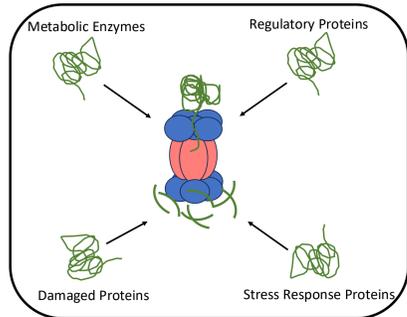
COLLEGE OF
SCIENCE & ENGINEERING

DEPARTMENT OF BIOLOGY

Aeron Pennington, Salina Hona, Josey Austin, Kelsey Waite and Shauna M. McGillivray
Department of Biology, Texas Christian University, Fort Worth, TX

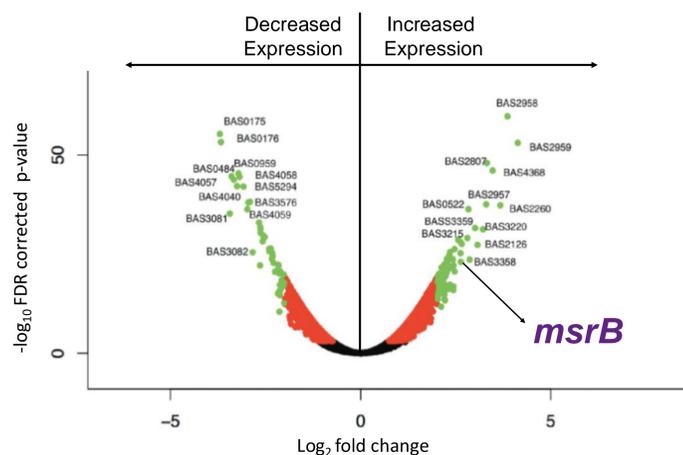
Background

ClpX is essential for virulence in *Bacillus anthracis* and critical for resistance to cell envelope-targeting antimicrobials. However, the exact mechanism behind this phenomenon is not yet fully understood. ClpX is a regulatory subunit of a major global protease, ClpXP.



ClpX: recognizes and unfolds proteins
ClpP: degrades proteins

Differential Gene Expression in $\Delta clpX$

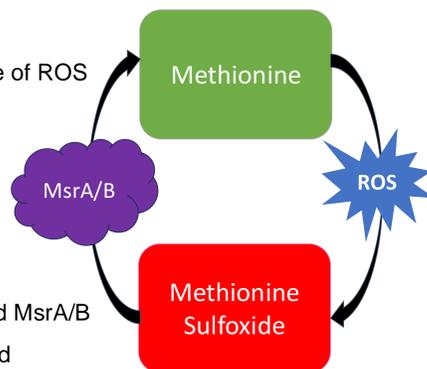


Claunch et al., 2018

Methionine Sulfoxide Reductase (MsrA and MsrB)

Methionine Sulfoxide Reductase:

- Antioxidant enzyme; linked to the tolerance of ROS
- Can exist independently or fused



In *B. anthracis*:

- No previous connection between ClpX and MsrA/B
- The role of MsrB has not been investigated

Fused MsrA/B:

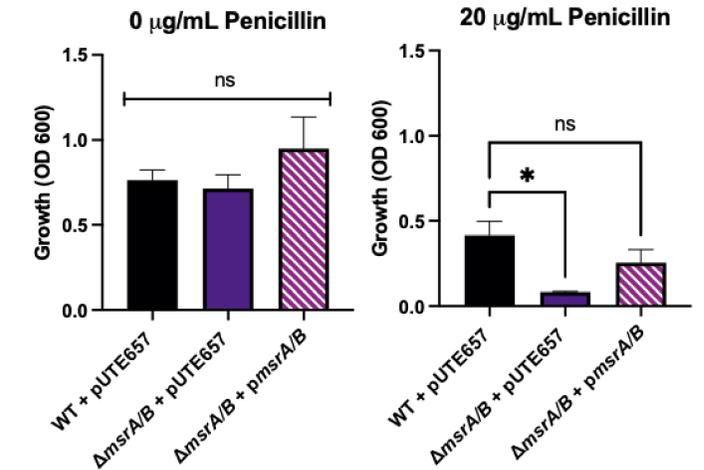
- At least 30 species of bacteria possess a fused *msrA/B*
 - Notably: *Streptococcus pneumoniae*, *Helicobacter pylori* and *Neisseria* spp.
- Fused MsrA/B possess similar enzymatic kinetics are independent units
 - In some species the MsrB subunit is more enzymatically active

Antibiotic MIC Table

Strains	Penicillin (μg/mL)	Daptomycin (μg/mL)	LL-37 (μg/mL)
WT	80	16	2
$\Delta clpX$	1.25	8	1
$\Delta msrA/B$	20	16	2

Cell wall Cell membrane

Complementation with $\Delta msrA/B$

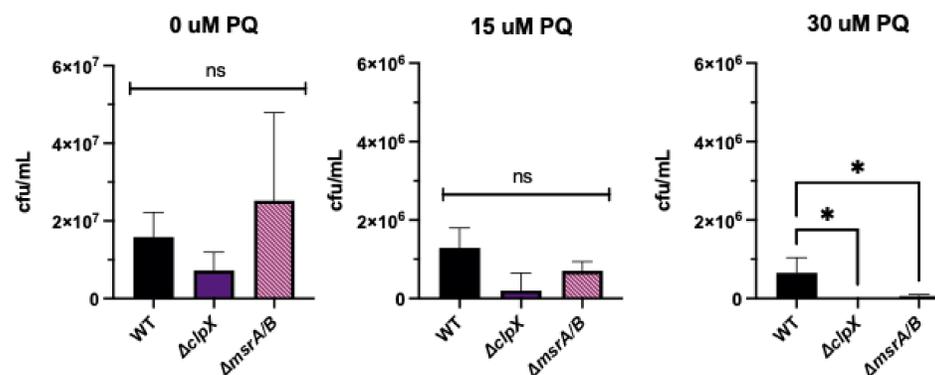


Data is represented as mean +/- SD of at least four independent experiments. ns indicates a non-significant difference, and * represents a p-value < 0.05, as determined by one-way ANOVA, followed by Tukey-Kramer *post hoc* analysis.

Susceptibility to ROS

ROS	H ₂ O ₂ (%)	Bleach (%)	Paraquat (μM)
WT	0.008	0.15	60
$\Delta clpX$	0.004	0.15	30
$\Delta msrA/B$	0.008	0.15	30

Survival in Paraquat (PQ)

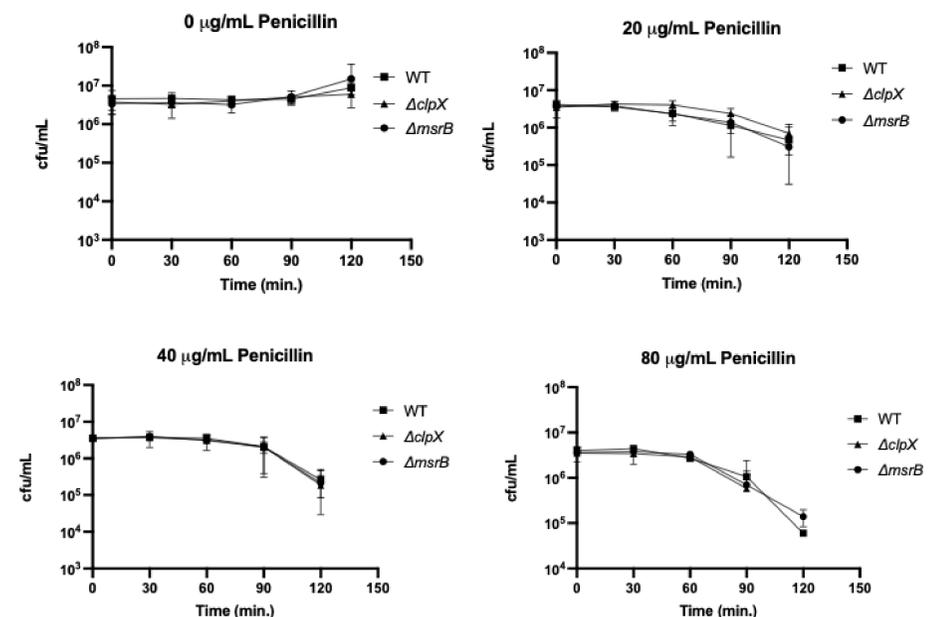


Data is represented as mean +/- SD of at least four independent experiments. ns indicates a non-significant difference, and * represents a p-value < 0.05, as determined by one-way ANOVA, followed by Tukey-Kramer *post hoc* analysis.

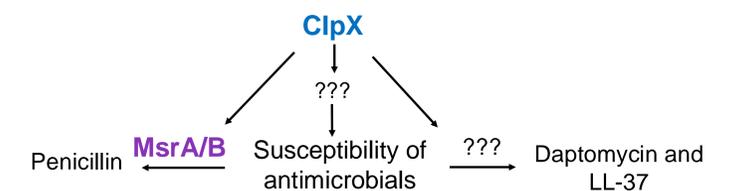
Future Directions

- Examine the regulation of *msrA/B* expression in WT and $\Delta clpX$ with and without penicillin and daptomycin
- Construct a complete knockout of *msrA/B* in WT *B. anthracis* Sterne
- Mutate critical cysteines in the active sites of the fused *msrA/B* for future assays

Survival Curve



Summary



Acknowledgments

Funding for this project is from the TCU Department of Biology and SERC. Thank you to all the members of the McGillivray lab for all their help and support with this project.