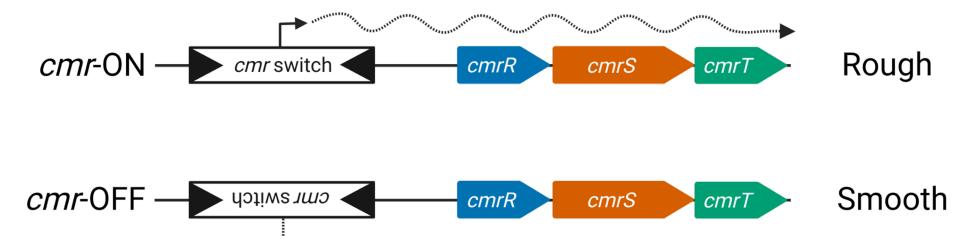
Identifying genes that affect colony morphology in bacterial pathogen *Clostridioides difficile*

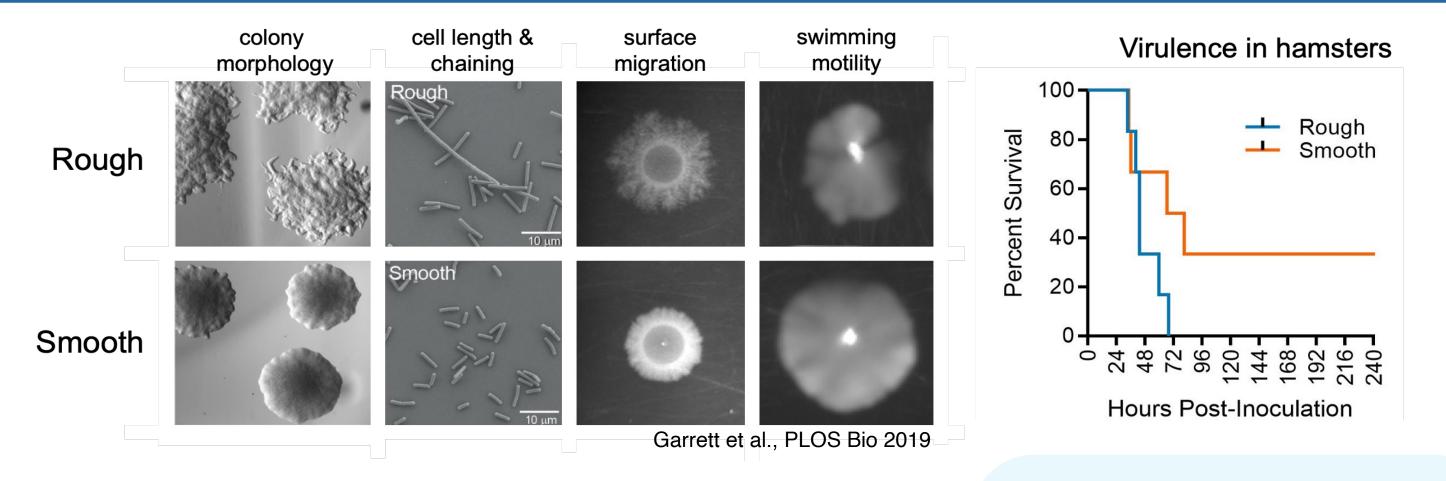
Aicha Kaouss, Anchal Mehra, Rita Tamayo, PhD

Department of Microbiology & Immunology, University of North Carolina at Chapel Hill

Phenotypic heterogeneity and virulence are linked in *C. difficile*

- *C. difficile* is a leading cause of healthcare-associated infections, and community-acquired cases are rising.
- Symptoms range from mild diarrhea to fatal pseudomembranous colitis and toxic megacolon.
- C. difficile exhibits phenotypic heterogeneity through phase variation a mechanism in which a bacterium can reversibly switch expression of specific genes on and off.
- Phase variation serves as a bet-hedging strategy to improve odds of survival in the presence of a changing or stressful environment.
- The *cmr* switch, one of eight known phase variable elements in *C*. *difficile*, modulates expression of *cmrRST*, which encodes an atypical signal transduction system.





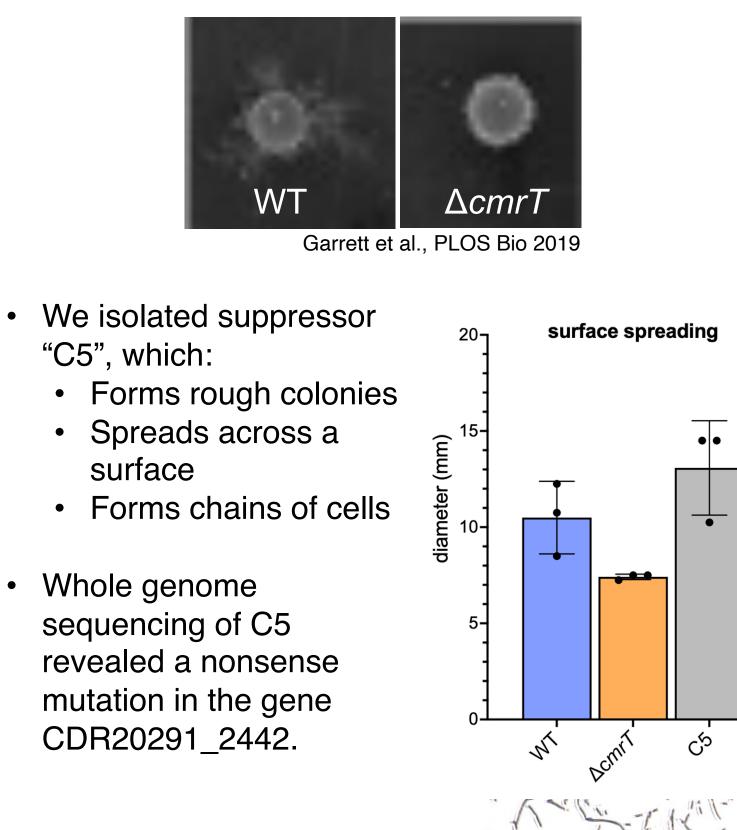
- Expression of *cmrRST* results in the formation of rough colonies, cell chaining, decreased swimming motility, increased surface motility, biofilm formation, and increased virulence in the hamster model of infection.
- Our group is taking multiple approaches to identify the genes that affect these phenotypes: RNA-Seq, suppressor screens, and transposon mutagenesis

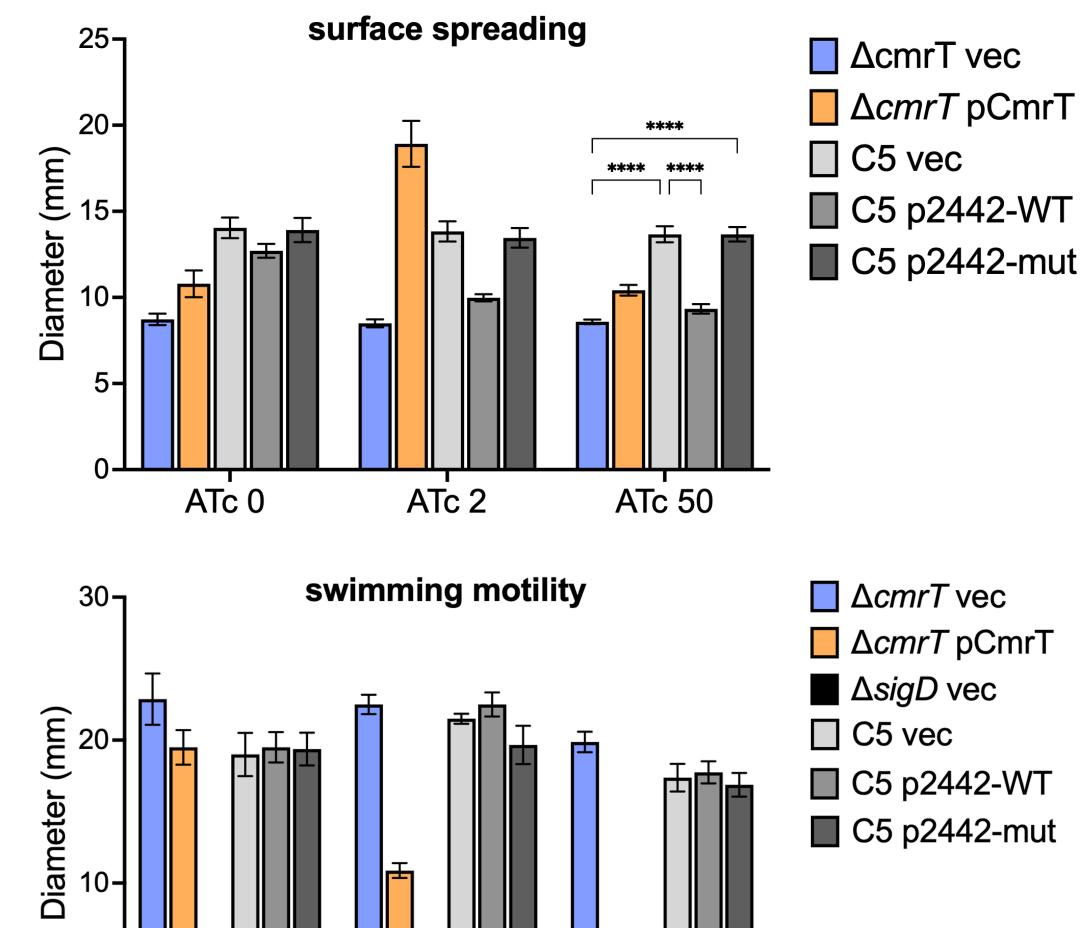
Research Goal: Identify genes important for rough or smooth colony formation via:

1. Suppressor Screen 2. Transposon Mutagenesis

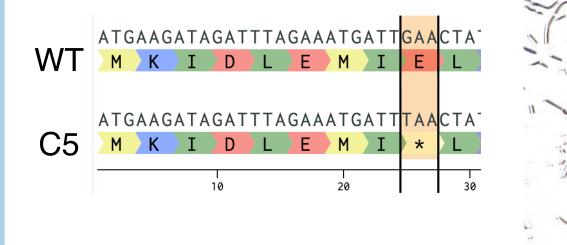
A suppressor mutation restores rough colony morphology in the $\Delta cmrT$ strain

• Using the $\Delta cmrT$ strain, which cannot form rough colonies or spread across an agar surface, we screened for suppressors that restored rough colony morphology and surface spreading after extended incubation on agar.

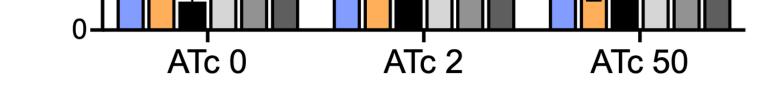




- $\Box \Delta cmrT$ vec $\Box \Delta cmrT pCmrT$ $\Delta sigD$ vec C5 vec C5 p2442-WT C5 p2442-mut
- ΔTv **Δ**T pT C5 pMut C5 pWT C5 v
- Complementing a functional, wild-type version of CDR20291_2442 into C5 restores $\Delta cmrT$ phenotype.
- *cmr*-ON-associated surface spreading correlates with decreased swimming motility. However, a swimming motility assay in the C5 suppressor revealed no significant difference in swimming motility between C5 suppressor and $\Delta cmrT.$







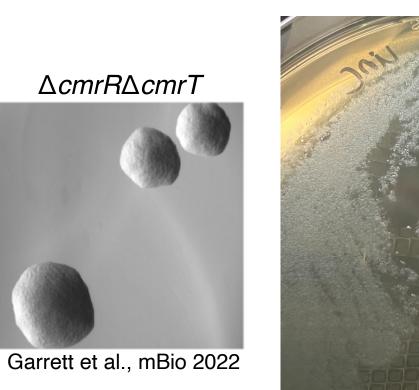
Wild-type CDR20291_2442 may function to maintain smooth colony morphology

 These findings suggest that swimming motility is regulated by a separate mechanism than surface motility in the C5 suppressor.

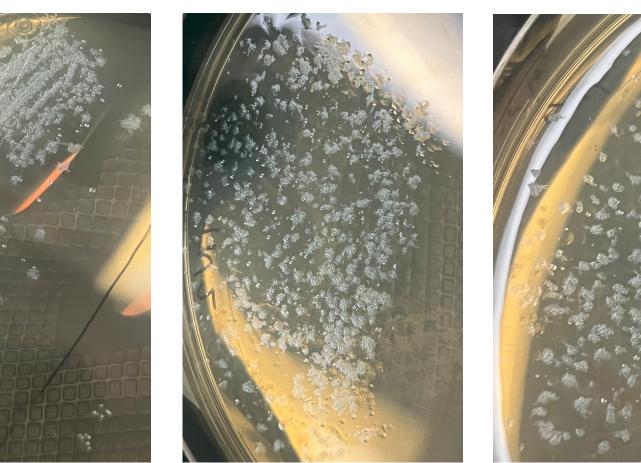
Using transposon mutagenesis to find isolates that restore rough colony morphology

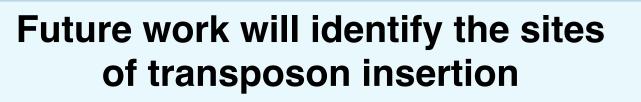
Approach:

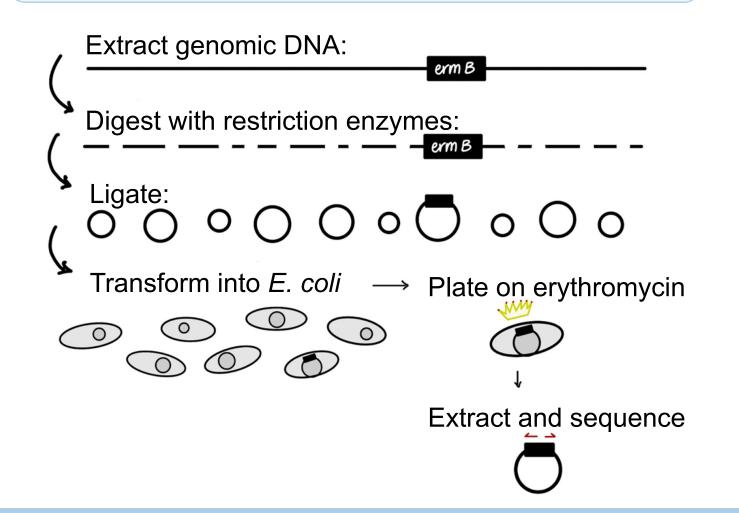
- 1. Plasmid carrying transposase and transposon inserted in $\Delta cmrR\Delta cmrT$ strain (forms smooth colonies)
- Transposition induced 2.
- Screen for rough colonies 3.
- 4. Rough colony isolates plated on antibiotics to ensure transposon insertion into genome



We identified ~30 colony isolates that displayed rough colony morphology.







NIH

RENDER

References

- 1. Garrett EM, Sekulovic O, Wetzel D, Jones JB, Edwards AN, Vargas-Cuebas G, et al. (2019) Phase variation of a signal transduction system controls Clostridioides difficile colony morphology, motility, and virulence. PLoS Biol 17(10): e3000379.
- 2. Garrett EM, Mehra A, Sekulovic O, Tamayo R. Multiple Regulatory Mechanisms Control the Production of CmrRST, an Atypical

Acknowledgements

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