

# Recent genetic drift in the co-diversified gut bacterial symbionts of laboratory mice



Daniel D. Sprockett<sup>1</sup>, Brian A. Dillard<sup>1</sup>, Abigail A. Landers<sup>1</sup>, Jon G. Sanders<sup>1</sup>, Andrew H. Moeller<sup>1,2</sup>

- 1. Department of Ecology and Evolutionary Biology, Cornell University
- 2. Department of Ecology and Evolutionary Biology, Princeton University

Co-diversified clades

ancestral to murids

#### Introduction

What are the evolutionary origins of the house mouse microbiota?





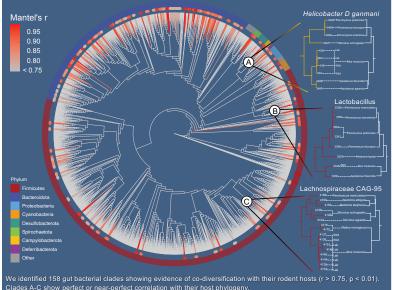
wild-living conspecifics, but we don't know if they've acquired those differences from the lab environment, or if lab lineages

#### Methods

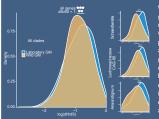


#### Results

Widespread co-diversification between gut bacterial symbionts and their rodent hosts

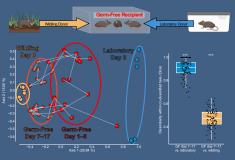


Significantly elevated genetic drift in lab strains



dN/dS in lab strains relative to dN/dS in individual genes car

Co-diversified wild strains out-compete related lab strains in vivo

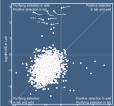


increased genetic drift, we re-analyzed data from competition experiments in germ-free mice<sup>2</sup>. Wild microbiotas were significantly

## Conclusions

co-diversified with rodents, and that these elevated levels of genetic drift during the last

# Altered genomic signatures of positive selection in lab strains



ancestral clades, we calculated derived MAGs. While most genes

## References

- Bowerman et al. (2021) Effects of laboratory domestication on the rodent gut microbiome
- Rosshart et al. (2021) Laboratory mice born to wild mice have natural microbiota and mode human immune responses. Science. DOI: 10.1126/science.aaw43
- Images created with BioRender.com



Funding



Graduate Student & Postdoc Opportunities Available



Contact: daniel.sprockett@wakehealth.edu