



The effect of simulated warming ocean temperatures on the bacterial communities on the shells of healthy and epizootic shell-diseased American Lobster (*Homarus americanus*)

Suzanne L. Ishaq¹, Grace Lee², Sarah M. Turner¹, Jean MacRae¹, Heather Hamlin¹, and Deborah Bouchard¹

¹ University of Maine, Orono, Maine, U.S.

² Bowdoin College, Brunswick, Maine, U.S.

American lobster (*Homarus americanus*)

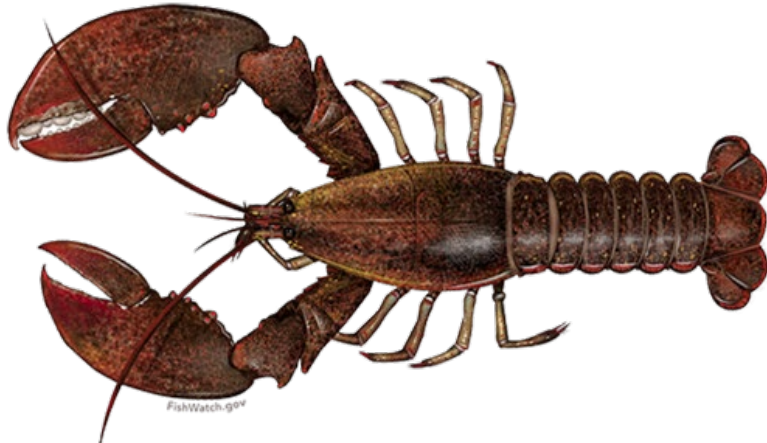


Image: <https://www.fisheries.noaa.gov/species/american-lobster>



Maine
pronunciation
guide: “lob-stah”

Image: <https://www.maine.gov/dmr/science-research/species/lobster/lobsterlicenseplate.html>

Marine crustaceans

Ecological impact not well studied in all lobster species

- Mid-trophic level: push energy from primary producers up the food chain)
- Scavenge detritus / turnover decaying material

Molt their shells to grow

Up to 100 year lifespan

Atlantic lobster industry

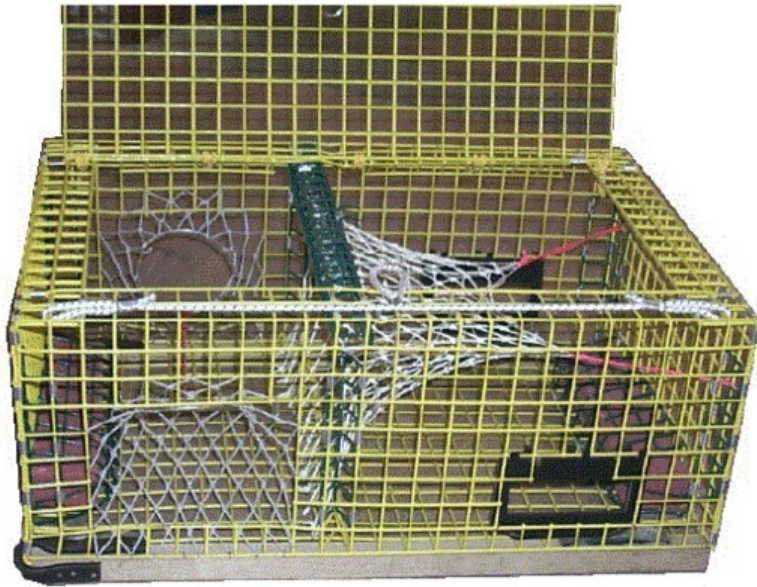


Image: <https://www.fisheries.noaa.gov/species/american-lobster#commercial>

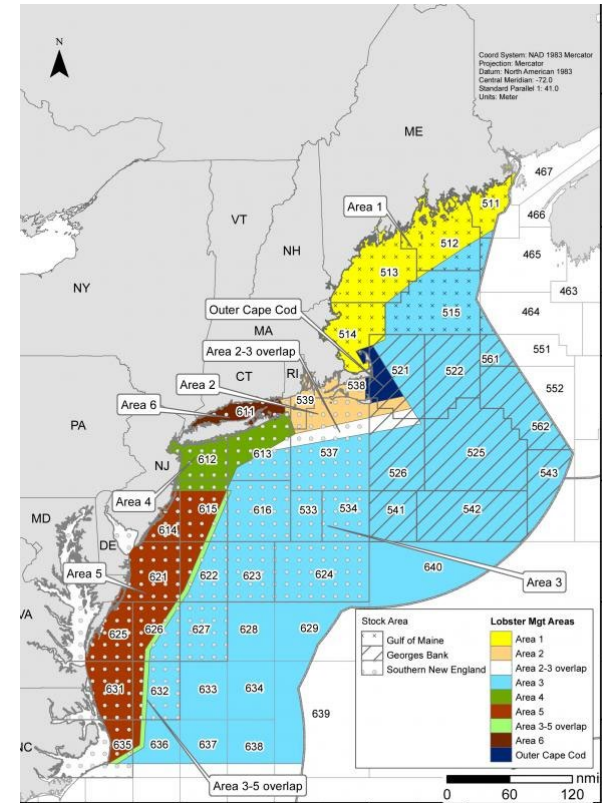
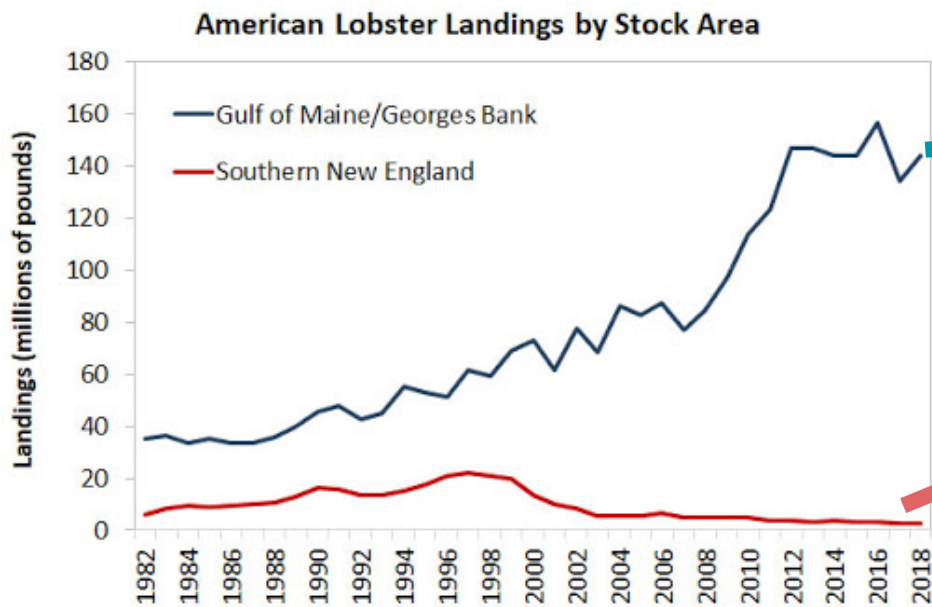


Image: penbaypilot.com/article/lobster-catch-maine-down-southern-new-england/57279

North Atlantic lobster industry



Source: Atlantic States Marine Fisheries Commission

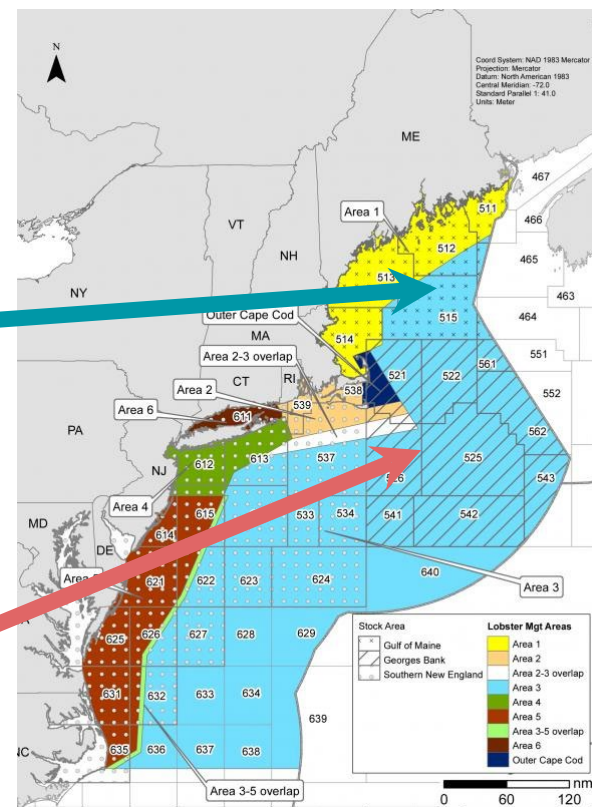
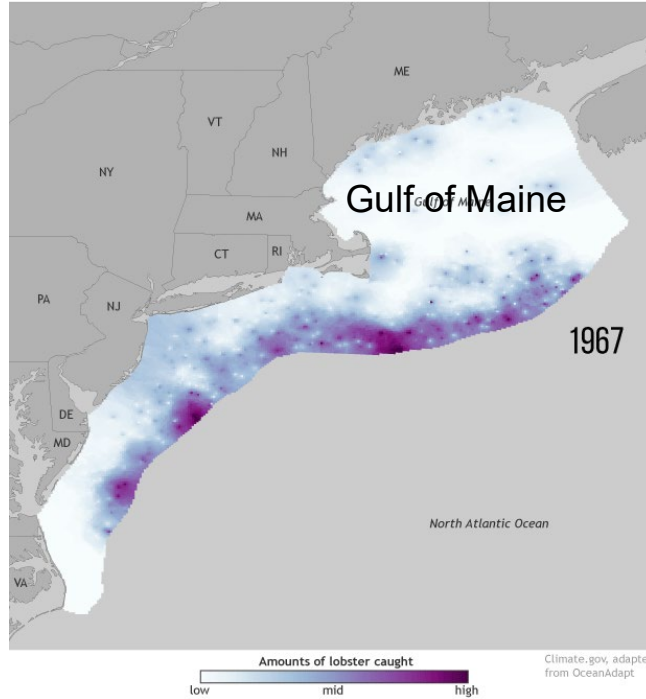


Image: penbaypilot.com/article/lobster-catch-maine-down-southern-new-england/57279

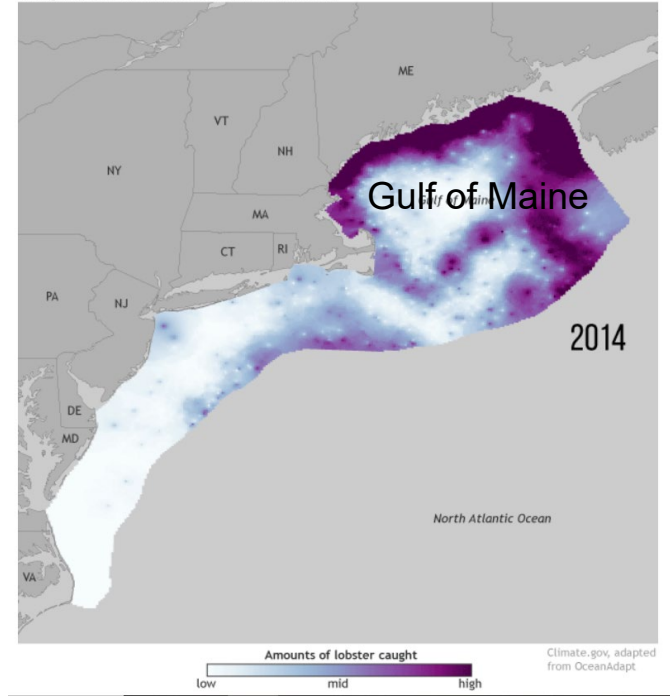


Warming coastal waters pushing lobsters north

Changes in lobster distribution from 1967 to 2014



Changes in lobster distribution from 1967 to 2014



Source: <https://www.climate.gov/news-features/climate-and/climate-lobsters>



Epizootic shell disease (ESD)

Causative agent(s) unknown

Causes pitting or softening of the shell

- Is not known to kill a lobster directly
- Increases susceptibility to infection, predation, physical damage

Advancing rapidly in wild populations

- Connection to ocean warming?

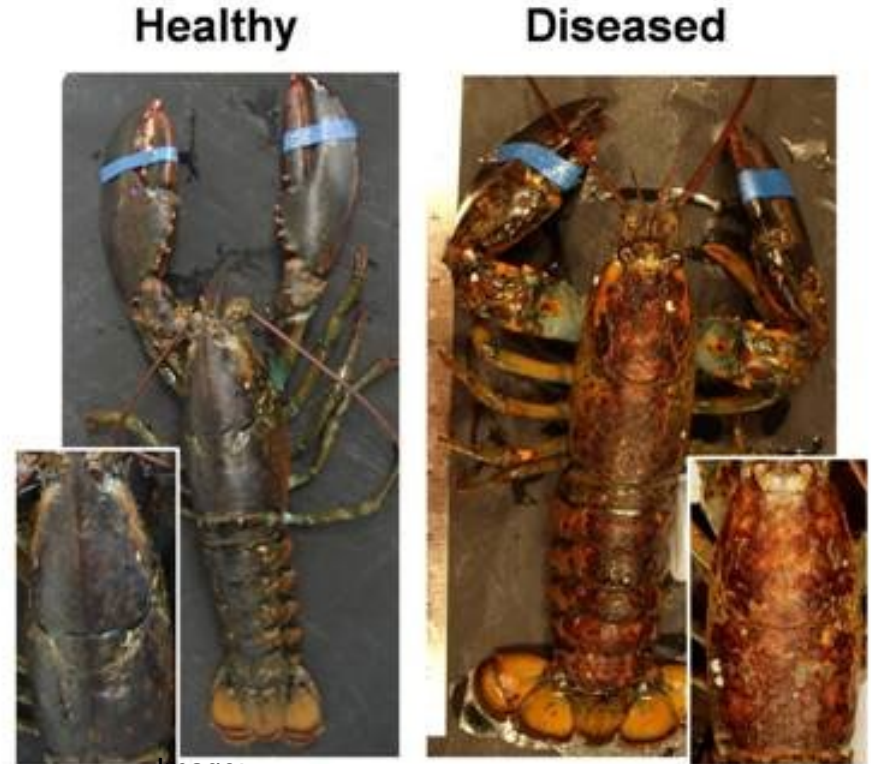


Image:

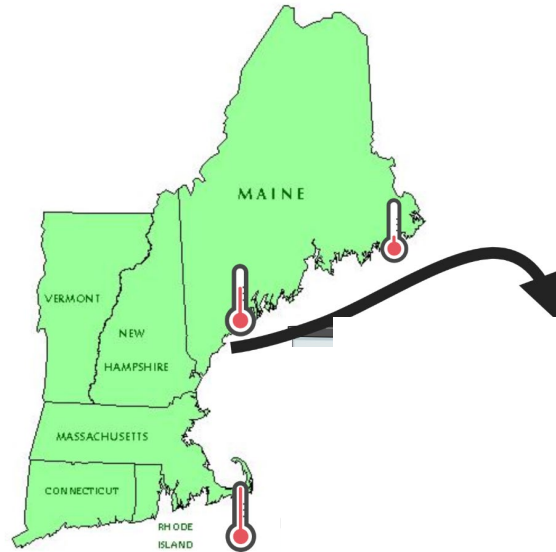
https://www.vims.edu/research/departments/eaah/programs/crustacean/research/lobster_shell_disease/



Experimental design

57 female lobsters collected from southern Maine coastal waters

Management zones F, G

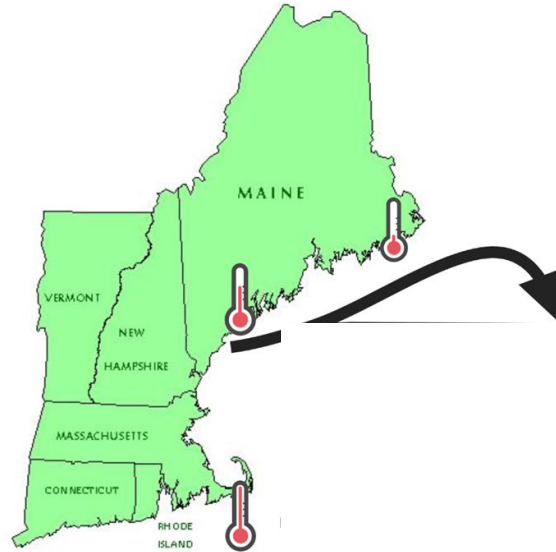


Created with BioRender.com

Experimental design

Lobsters were assessed for visible shell disease and scored by percent of shell covered (Smolowitz et al., 2005)

- 0, no observable signs
- 1, on 1 - 10% of shell
- 2, on 11 - 50% of shell
- 3, on >50% of shell



Created with BioRender.com

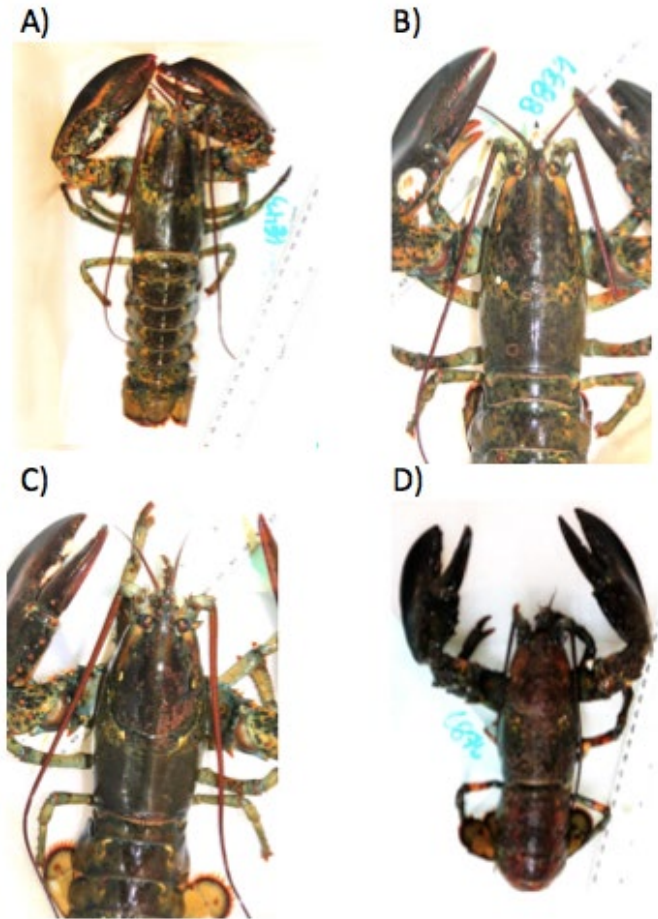


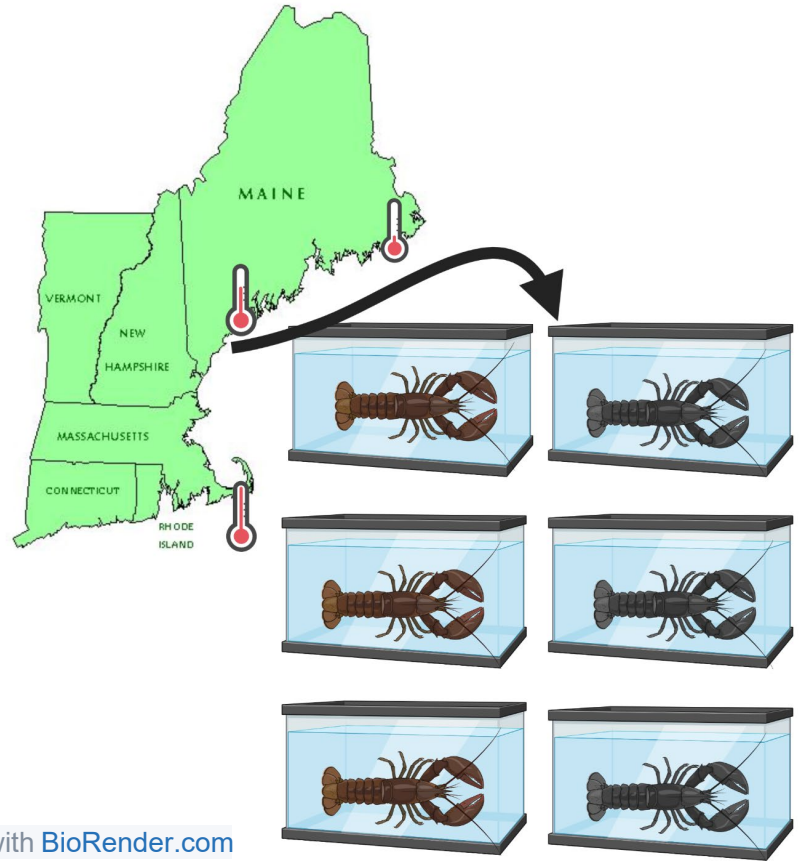
Image: Deborah Bouchard

Experimental design

Lobsters were placed into separate tank systems based on

- A) Apparently healthy
- B) Shell Disease Index 1 - 3

Lobsters housed in individual tanks



Created with BioRender.com

Experimental design

2 weeks of adaptation

Baseline sampling
(summer)

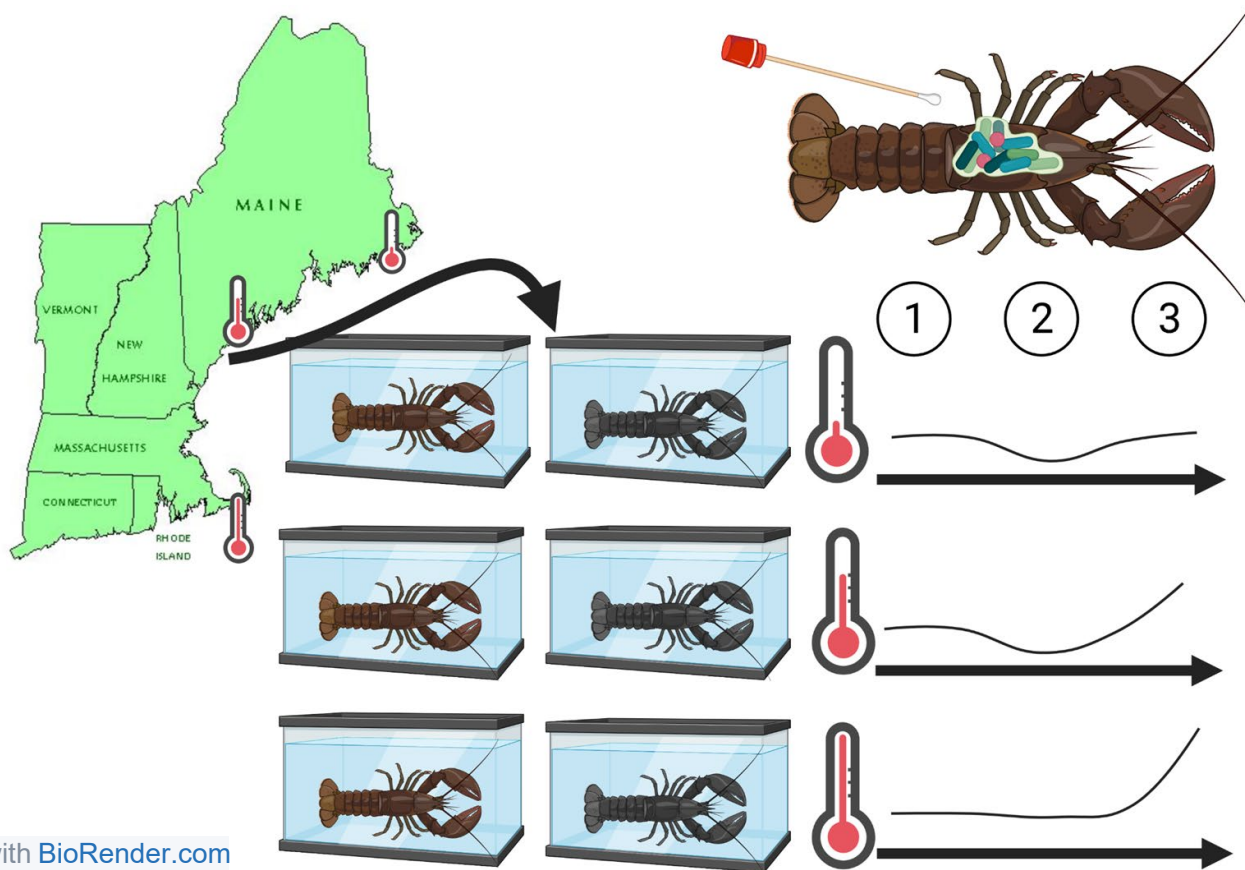
- Tank water a 11°C

4 months later (winter)

- Tank water respective to regional simulation

10 months later (summer)

- Tank water respective to regional simulation



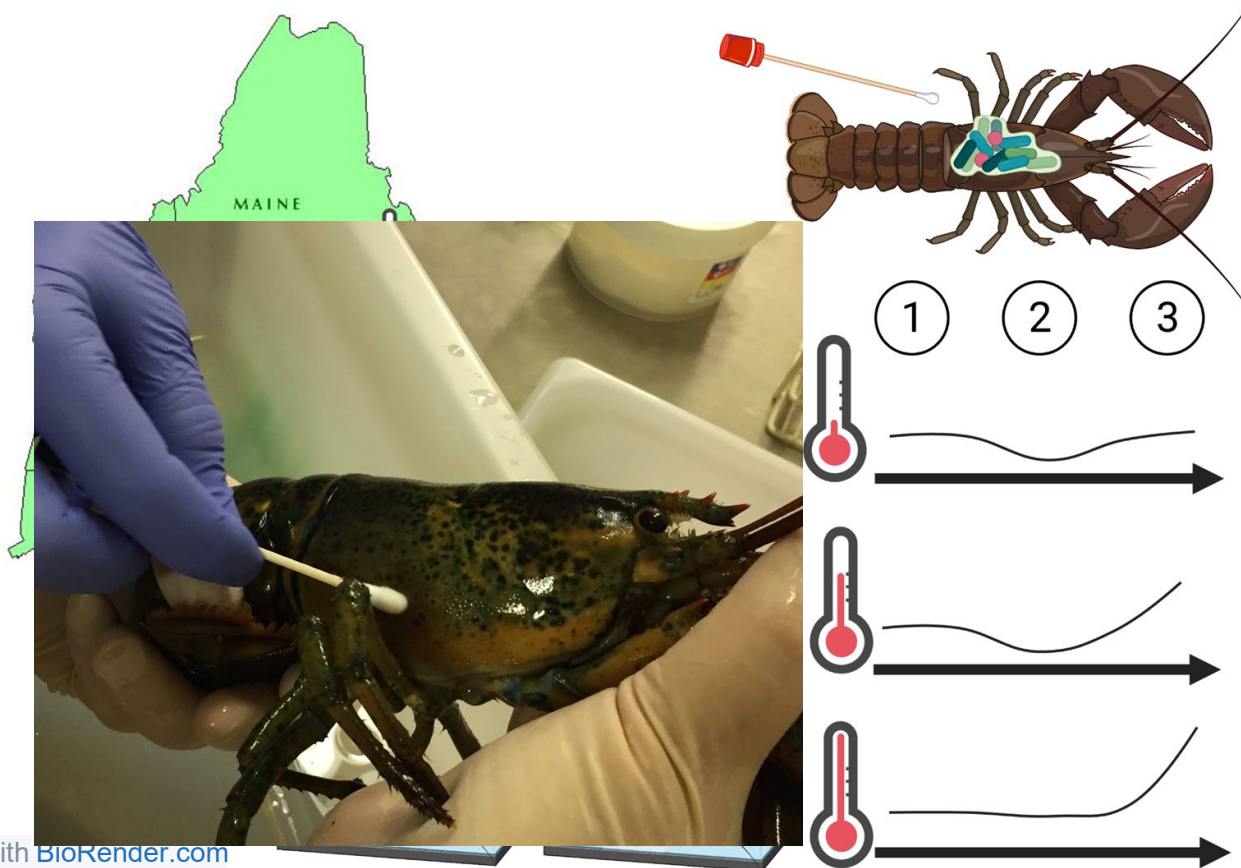
Created with BioRender.com

Experimental design

Samples collected at 3 timepoints

- Shell bacterial community, lobster weight, lobster length, Shell Disease Index

A designated portion of the shell was swabbed to sample shell-associated microbial community



Created with BioRender.com

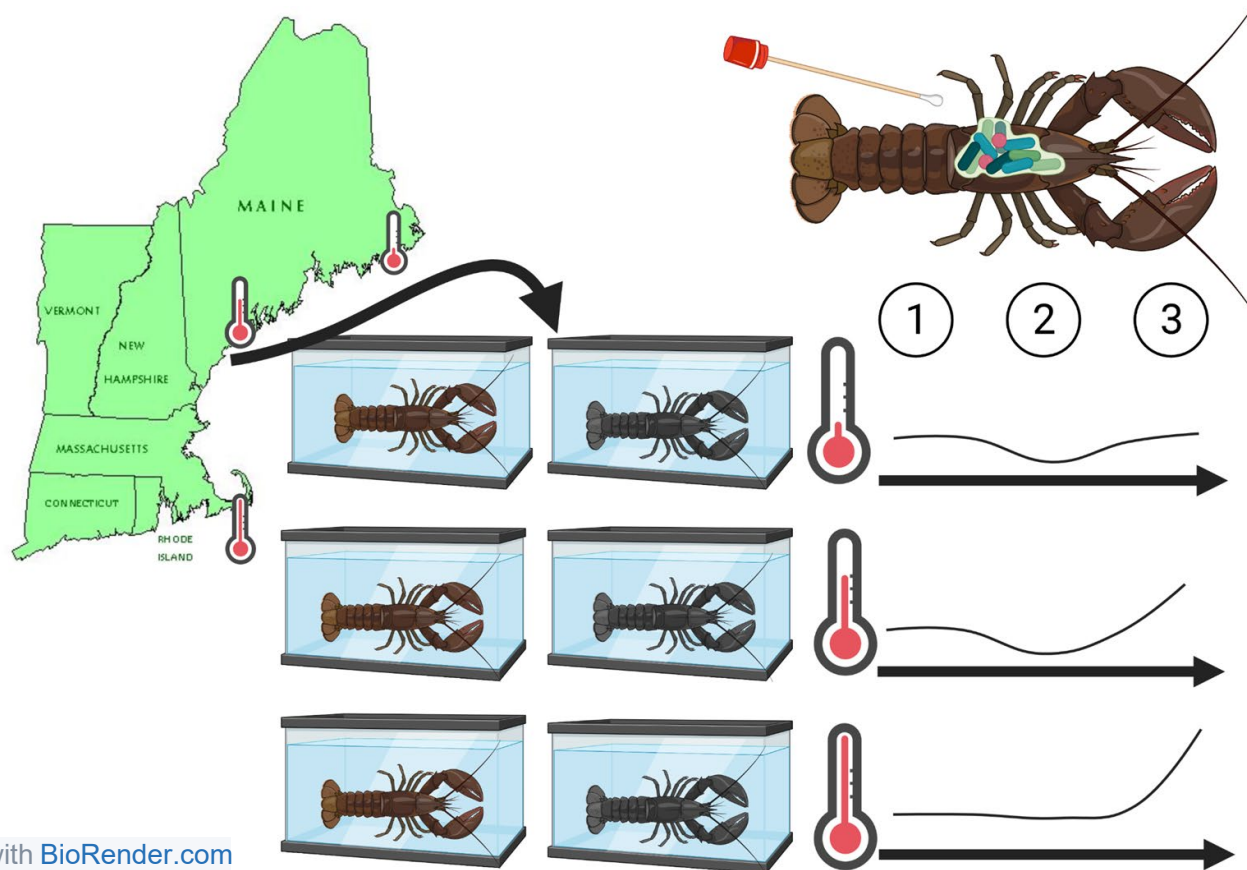
Experimental design

131 experimental samples, plus 10 controls

- PCR amplification and purification,
- Illumina MiSeq ver. 4 sequencing,
- passed quality-control filtering.

Sequences were processed using the R software platform, using DADA2, phyloseq, vegan, and assorted other packages.

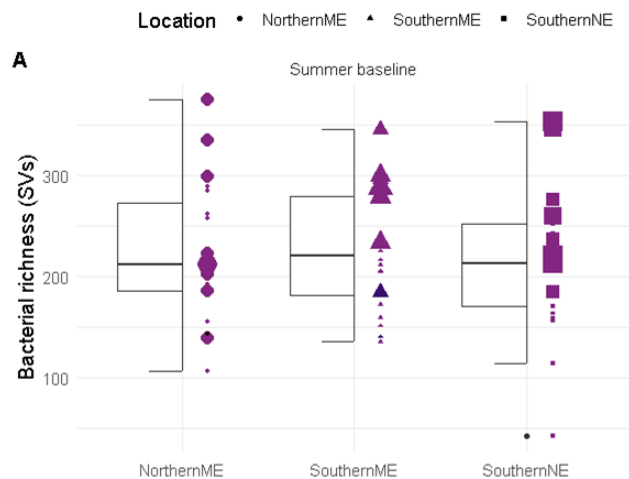
Created with BioRender.com





Bacterial richness on shells was same at baseline (as expected)

of bacterial taxa identified



All lobsters collected from same location

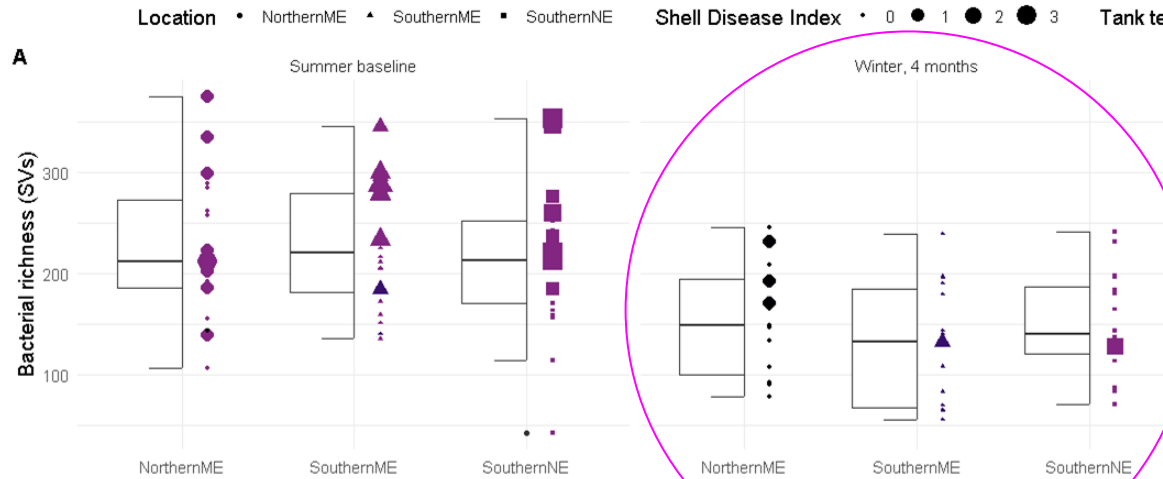
Baseline is 2 weeks after collection, all at 11°C

Point shape = location the tank temperature will mimic



Bacterial richness on shells affected by tank temperature

of bacterial taxa identified



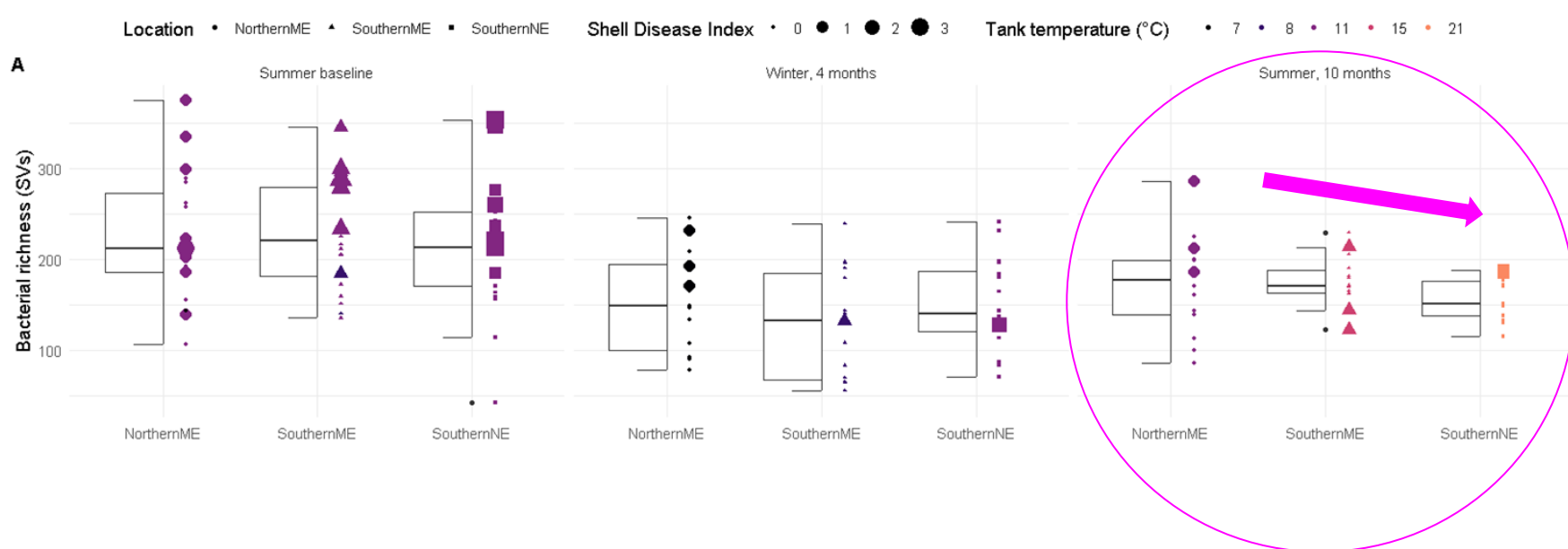
After 4 months in tanks, all lobster shells hosted fewer bacterial taxa

Starting to see differences by simulated geographic ocean temp



After 10 months, lobster shells in warmer water host fewer different bacterial taxa

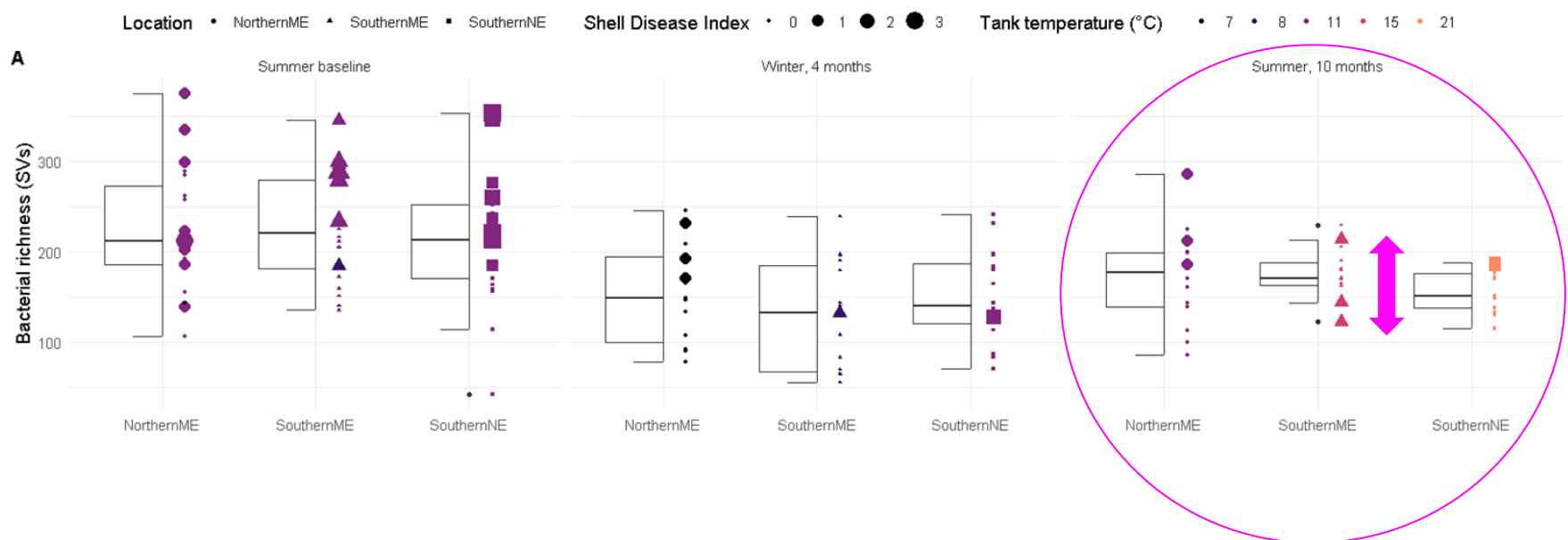
of bacterial taxa identified



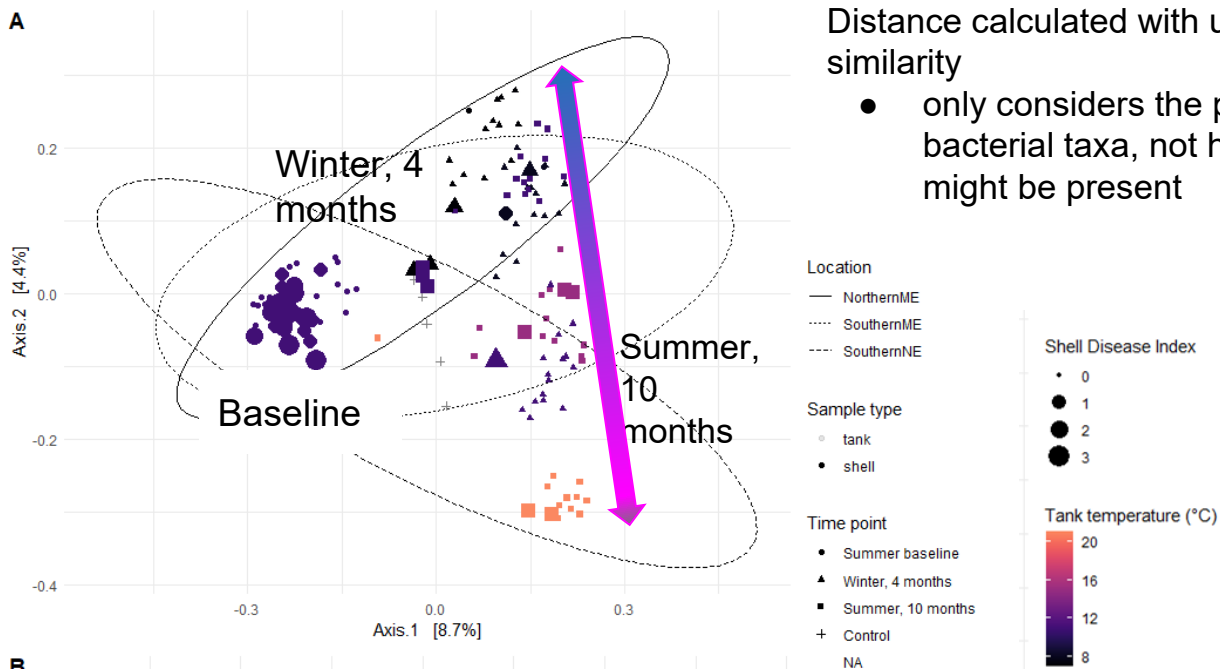


Lobster shell bacterial community richness in warmer water were less variable

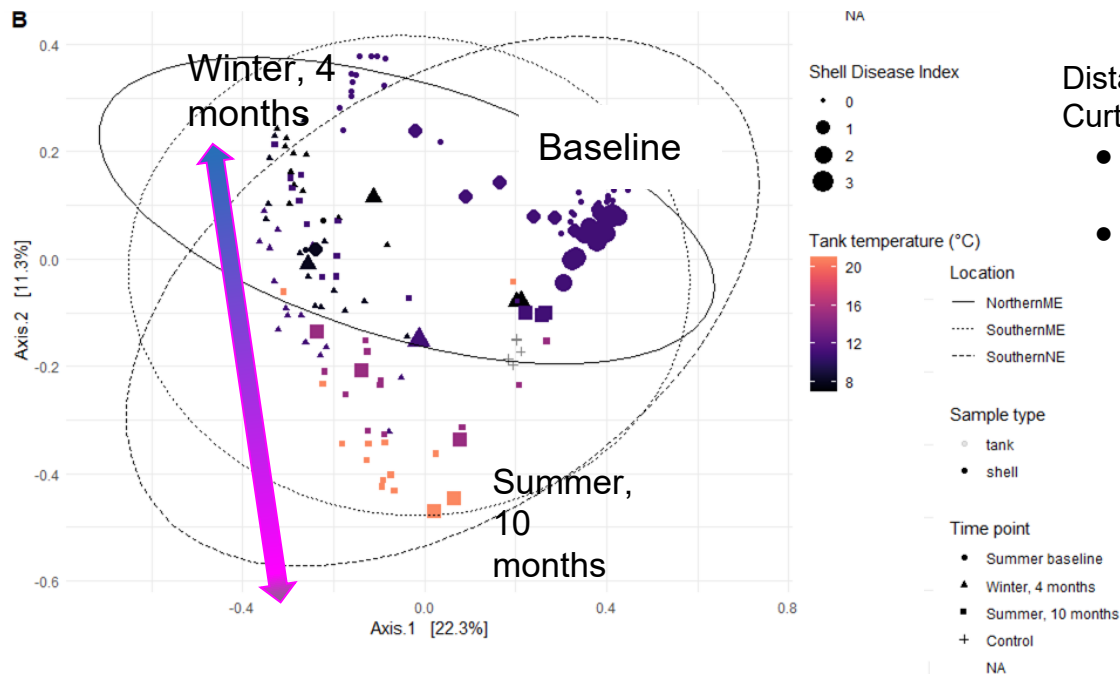
of bacterial taxa identified



Tank temperature affects “who” is found on lobster shells



Tank temperature affects abundance of each bacterial taxa found on lobster shells

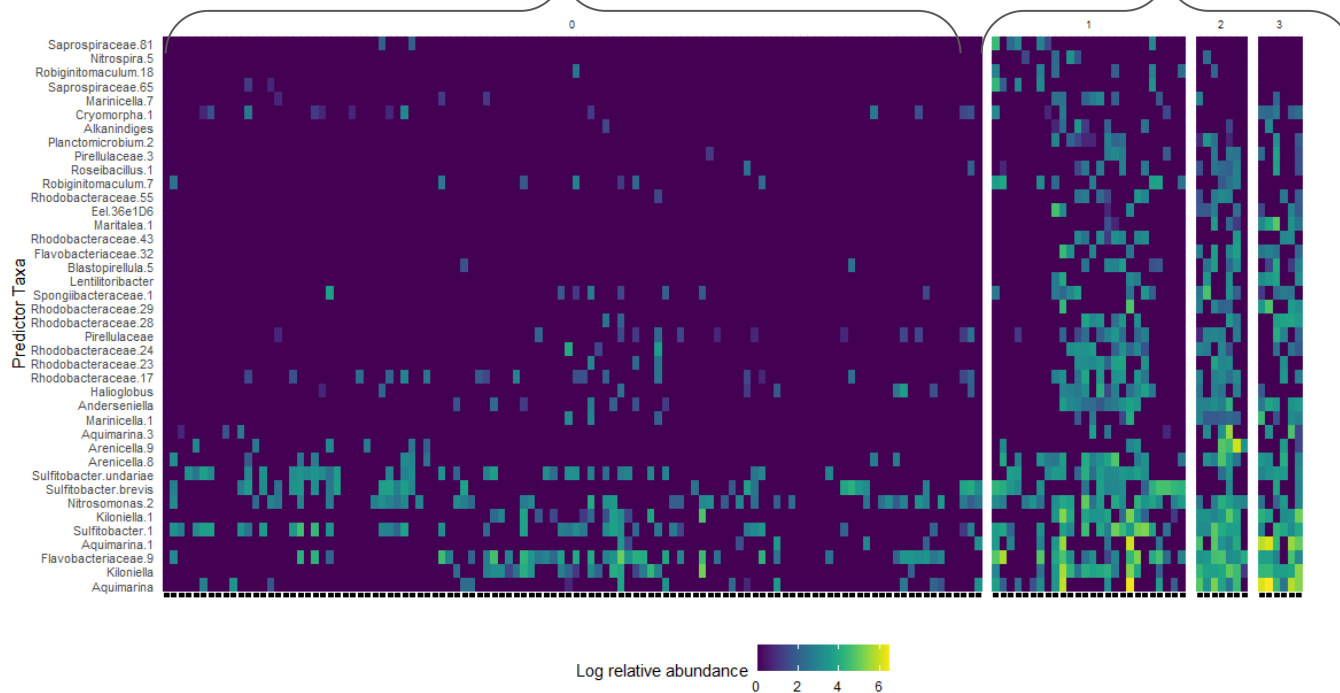


Distance calculated with weighted Bray-Curtis similarity

- considers the presence/absence of a bacterial taxa
- AND how many of that taxa might be present (abundance)



Subtle changes to the bacterial community in healthy vs. epizootic shell diseased



Broader impacts of this work

New England lobster industry survey:

- Been lobster fishing for decades
- Most have significant financial investments in their business (personal funds, loans)
- 9 - 36% have not finished high school
- In Maine especially, few local economic alternatives

Surveyed by Gulf of Maine Research Institute, 2008,
http://www.lobstermen.com/wp-content/uploads/2009/10/RES_DH_reports_Lobster-Socioec-Survey.pdf

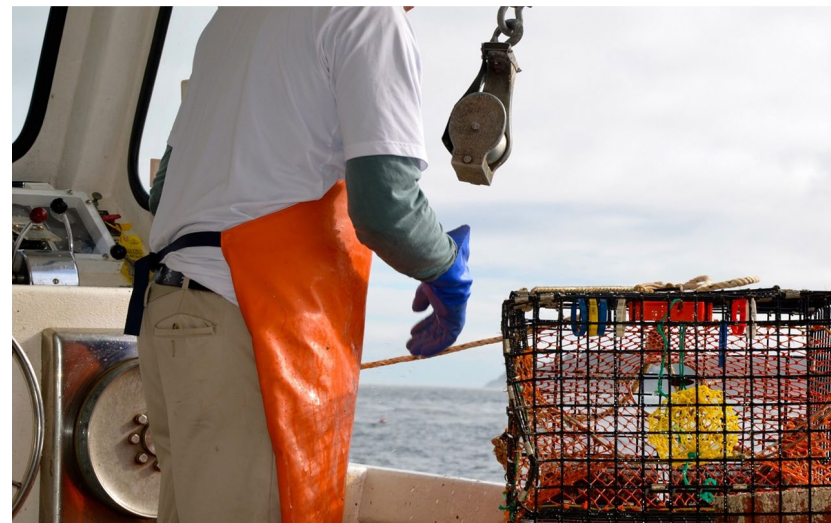


Image: <https://pixabay.com/photos/boating-lobster-fisherman-fishing-1343347/>



Acknowledgements

- State of Maine Department of Marine Resources for collecting lobsters for us
- Maine Lobster Research Education and Development (RED) board for funding the research (DMR ME CT#13A-20160606*3908)
- University of Maine Aquatic Animal Health Laboratory Staff for assisting in all phases of the research, M. Scarlett Tudor, Emily Thomas, Dawna Beane, as well as University of Maine undergraduate students Laurel Anderson, Emily Tarr, and Helen Reese