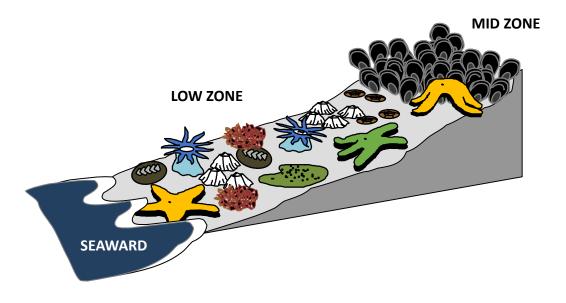
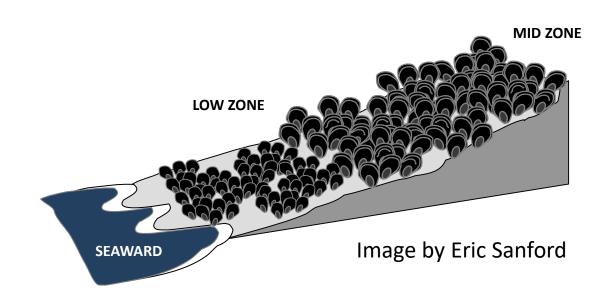


## Mechanisms of resistance, resilience to SSWS

### Pisaster present



#### Pisaster removed



- Recruitment of mussels and barnacles
- Large predators (stars and birds)

- Facilitation by algae, barnacles
- Small predators

Can compensatory predation by *Nucella ostrina* and *Leptasterias spp.* prevent downward spread of mussels into the low zone?

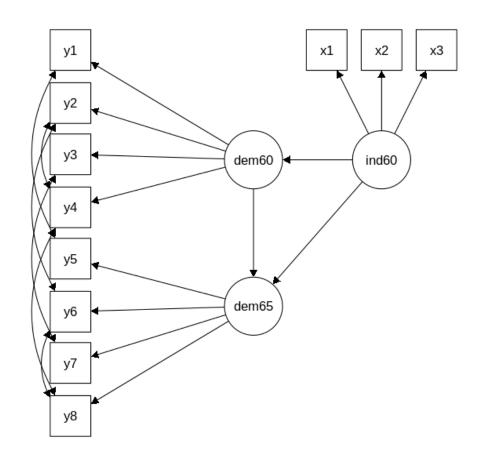
• How does the strength of compensatory predation vary from SoCal to Oregon?

 Hypothesis: higher densities of predator species in OR=increased compensatory predation

 How do recruitment and colonization dynamics vary from SoCal to Oregon over the course of ~16 months?

### Products of this class

- Work through vignettes in R package lavaan (and visualize with semPlot)
- Build an SEM for SPITFIRE- conceptual framework
- Stretch goal: use SP SEM to generate a dataset that, when model is fitted, returns the same parameter estimates used to generate the dataset

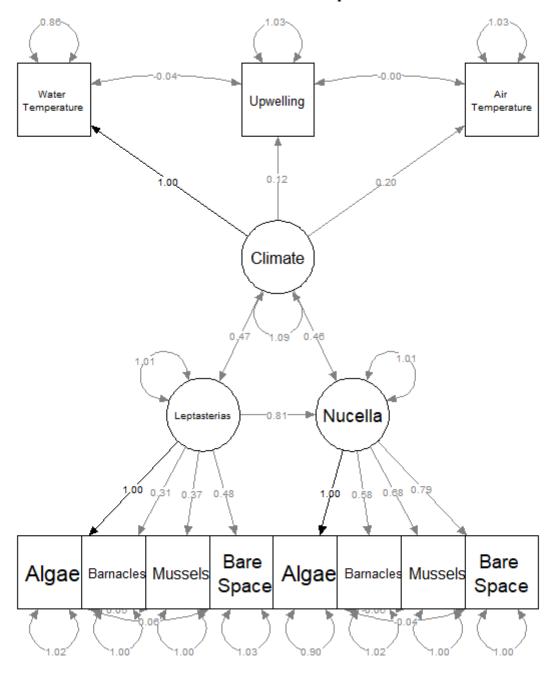


Can compensatory predation by *Nucella ostrina* and *Leptasterias spp.* prevent downward spread of mussels into the low zone?

• What is the relative strength of predation for each predator species on four functional groups of prey?

• Effects of environment (water temperature, air temperature, upwelling index) on predators

#### SPITFIRE structural equation model



# What was involved in getting to the model

Working through Lavaan vignettes

Reading/research- application of SEM in biological systems, how SEM works

 Troubleshooting model parameterization + fitting; drawing tons of conceptual models

## Workflow

• Part of workflow is currently missing: don't have dataset in hand

 Data simulation + model parameterization is one script; easily reproducible

 "troubleshooting" branch for testing data simulation functions; main branch has vignettes + models

# Ongoing work

1. Finish data processing

2. Assess model fit to my dataset

3. Make changes as needed- "model generating"

4. Profit



# Stretch goal

 Developing a theoretical SEM that's good enough to simulate data

 Can use to assess predictive capacity of the model once I have fit it to my actual data + ground truth mechanisms



#### Your population vs. your model



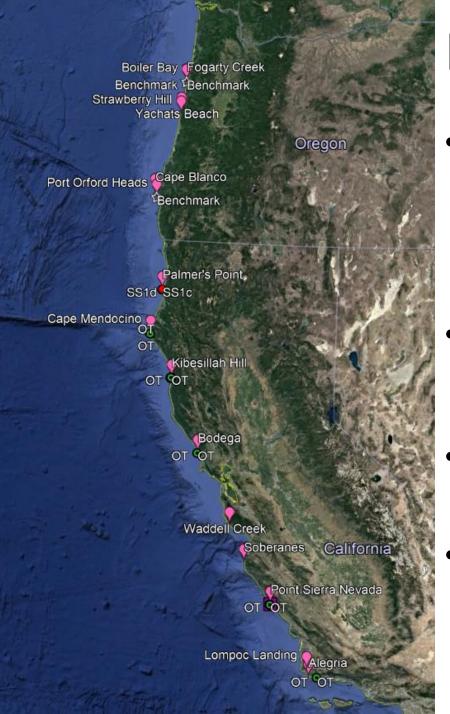
18:46 - 18. Nov. 2018

## Desired end result

Workflow for example previously described

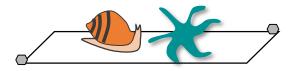
Drawn version of my actual model

 Goal is to have a flexible workflow that I can modify to run my data once I have it



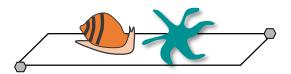
## Methods

- 2 plots, 1 fence, 4 cages per site
  - OR: Nucella only, Nucella and Leptasterias, Leptasterias only, no preds (control)
  - CA: Nucella only, no preds (control)
- 15x15 cm cages deployed ~1m under lower limit of mussel bed
- Cleared at experiment start (May 2018)
- Stocked w/preds (site specific density) once prey had recruited
  - OR/NorCal: July 2018
  - CenCal: September 2018 (low recruitment)



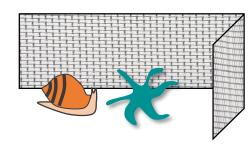
Recruitment plot (cleared ~monthly)

+ Nucella ostrina + Leptasterias



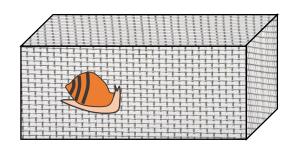
Colonization plot (cleared once)

+ Nucella + Leptasterias

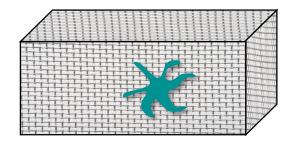


Colonization fence (Cage Control)

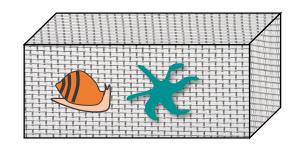
+ Nucella + Leptasterias



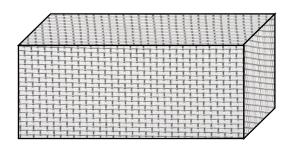
Nucella only + Nucella - Leptasterias



Leptasterias only
- Nucella + Leptasterias
OR only



Both predators + Nucella + Leptasterias OR only



Predator exclusion
- Nucella - Leptasterias

## Overarching goals- did I achieve them?

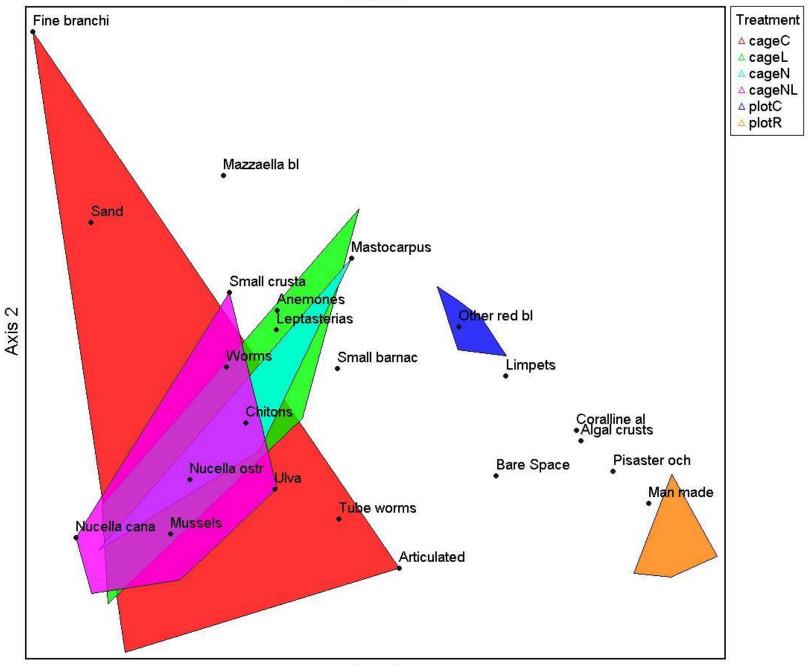
• Build a Structural Equation Model (SEM) – multivariate method that facilitates investigation of direct and indirect interactions

 Investigate mechanisms behind variation in interaction strengths across a wide biogeographic gradient

 Increasing understanding of mechanisms shaping community structure -> increased predictive capacity re: community-level response to climate change

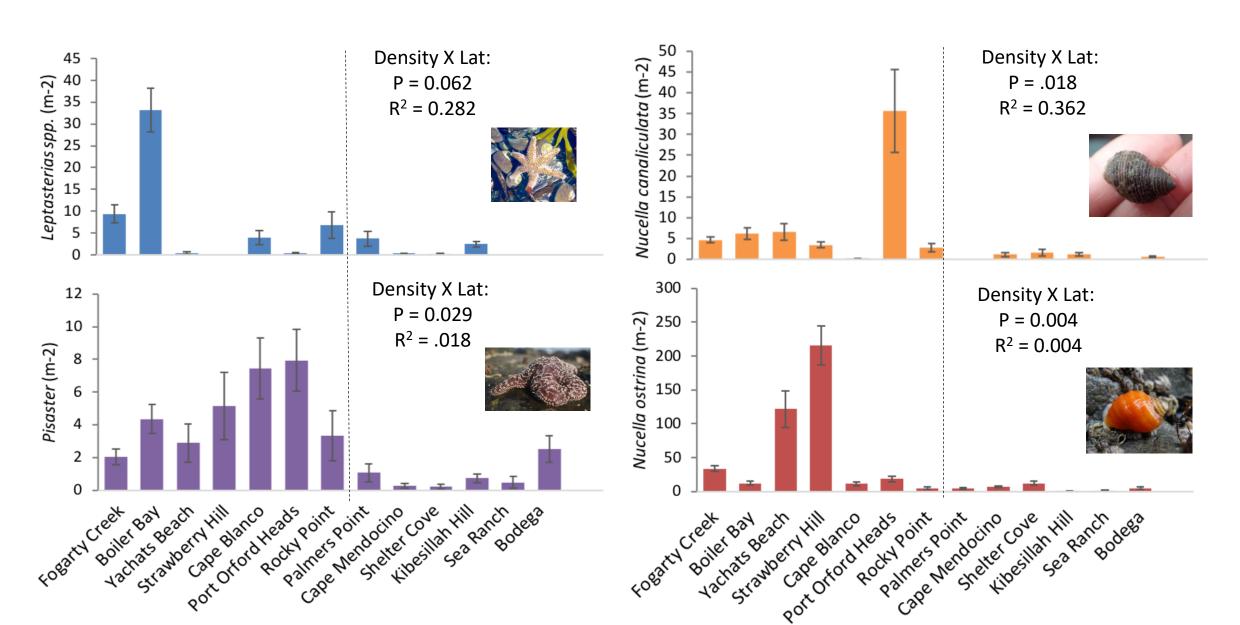


#### SH NMS



Axis 1

## **Predator Densities**



### The system is both spatially variable

### ...and temporally variable

