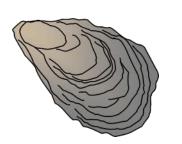
Oyster Model

Objective: evaluate potential impacts of future changes in hypoxia on living resources in Chesapeake Bay – oysters provide a test organism

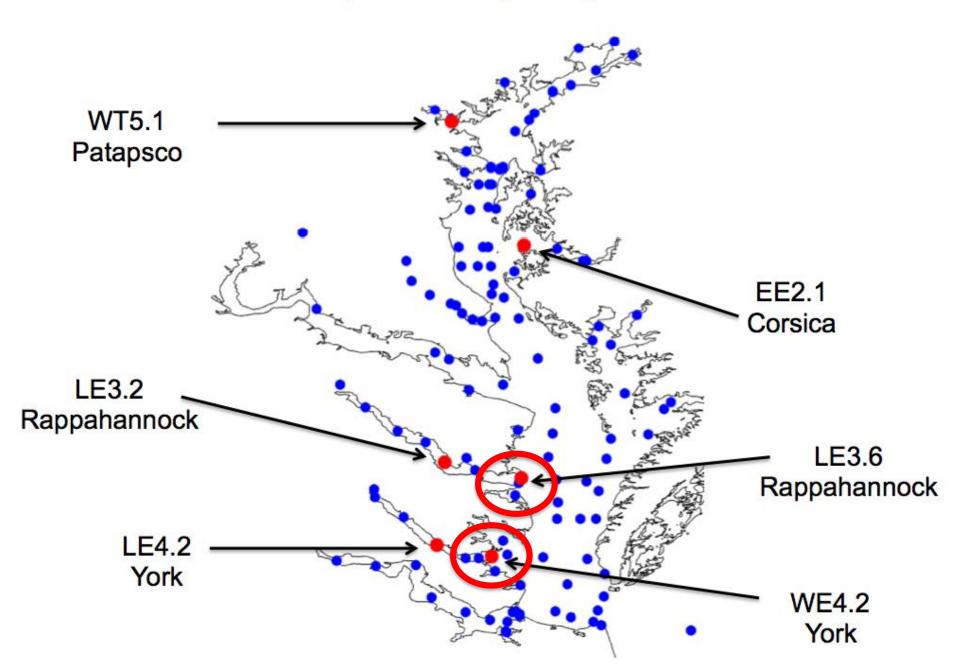


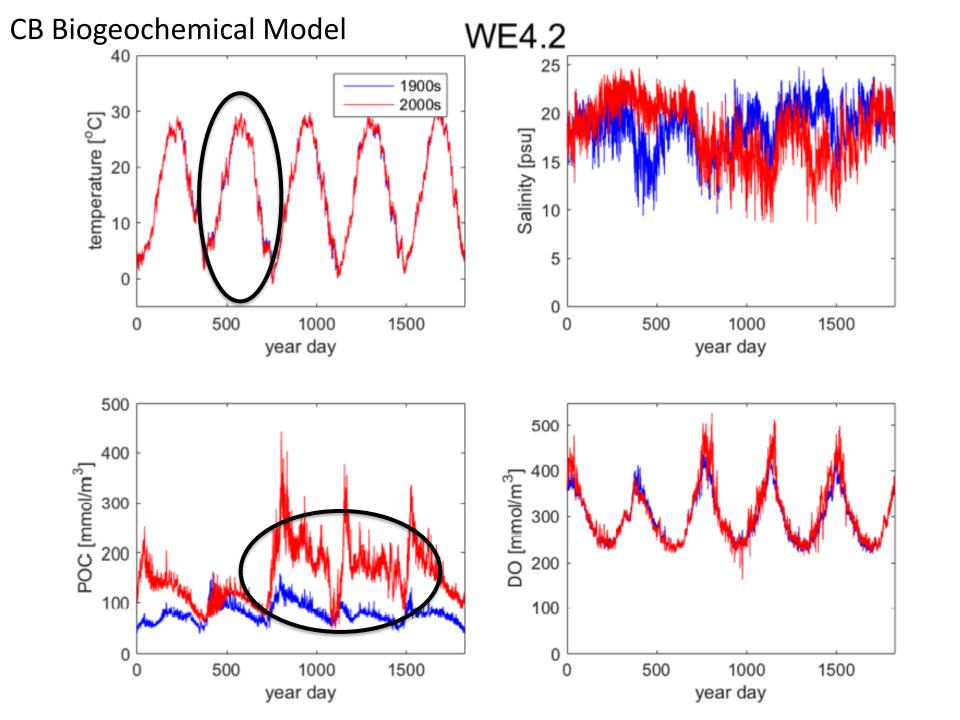


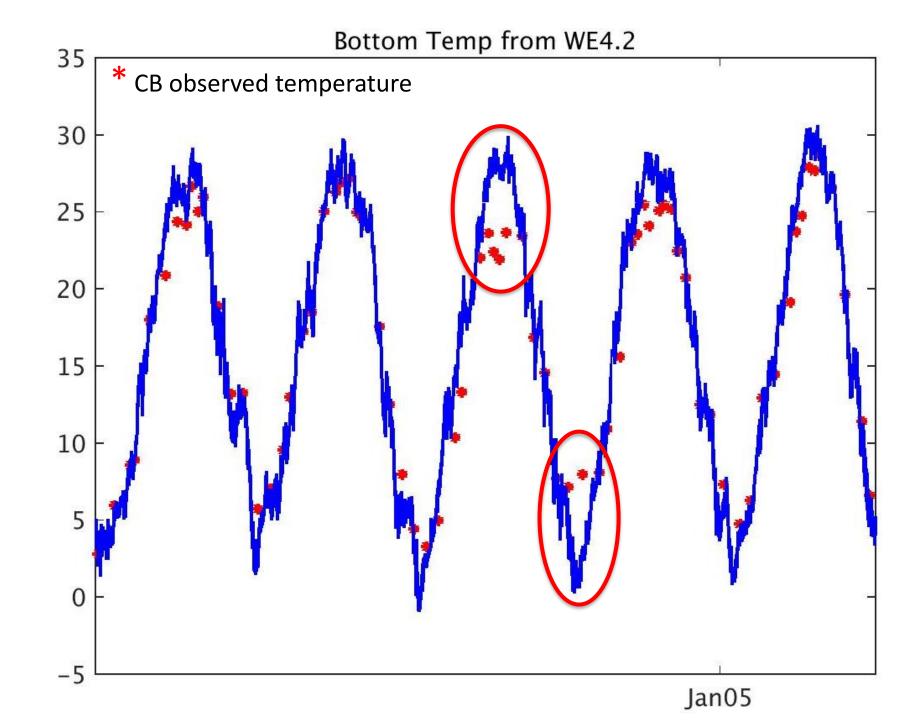
Oyster Model

- Focus is Crassostrea virginica (Eastern oyster)
- Model simulates post-settlement population dynamics (growth, reproduction, mortality)
- Includes larval, Dermo disease, MSX disease and genetics submodels
- Inputs are temperature, salinity, food, turbidity, bottom velocity
- Outputs are total number, population size frequency, reproductive capacity, mortality distribution, disease prevalence and intensity
- Model applied in a time-dependent mode at specific locations in Chesapeake Bay

Chesapeake Bay Program Stations



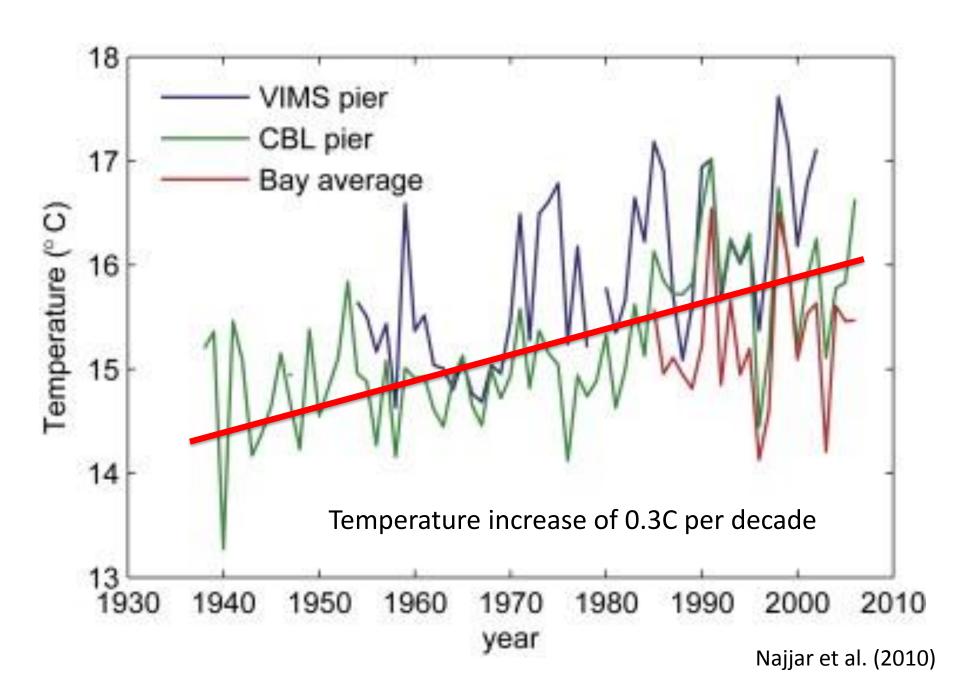




Oyster Model Forcing

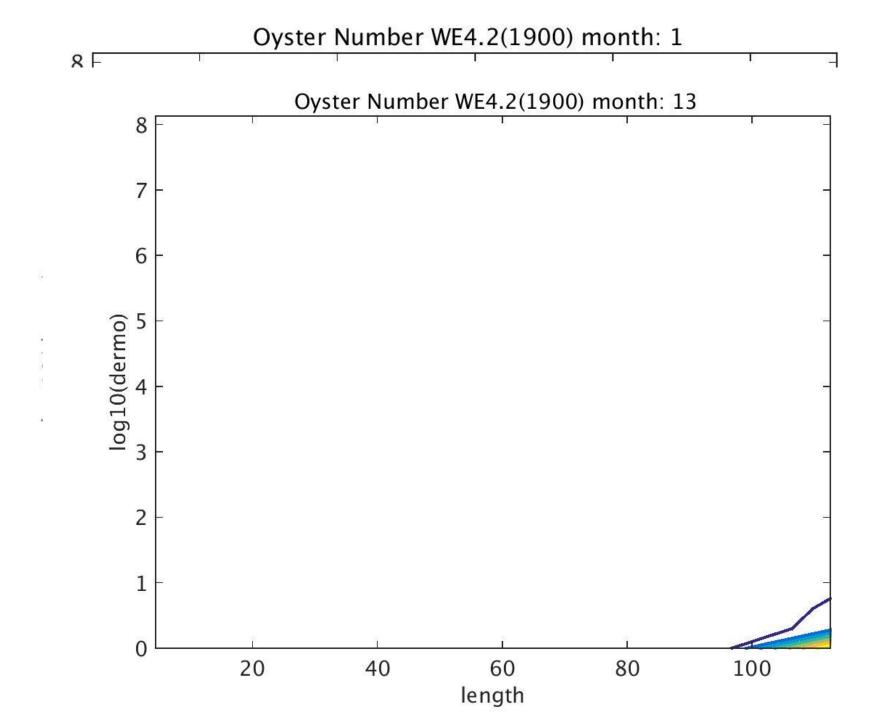
- Input salinity and velocity time series from CB simulations
- Used CB monitoring program bottom temperature for 2000-2005 for the oyster model
- Adjust CB monitoring program bottom temperature for 1900-1905
- Adjust food values for 2000-2005



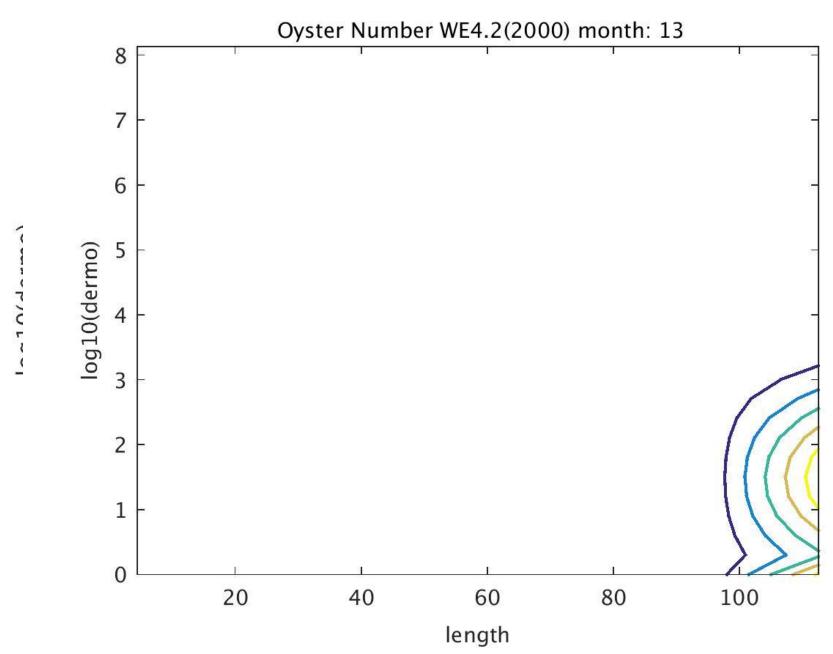


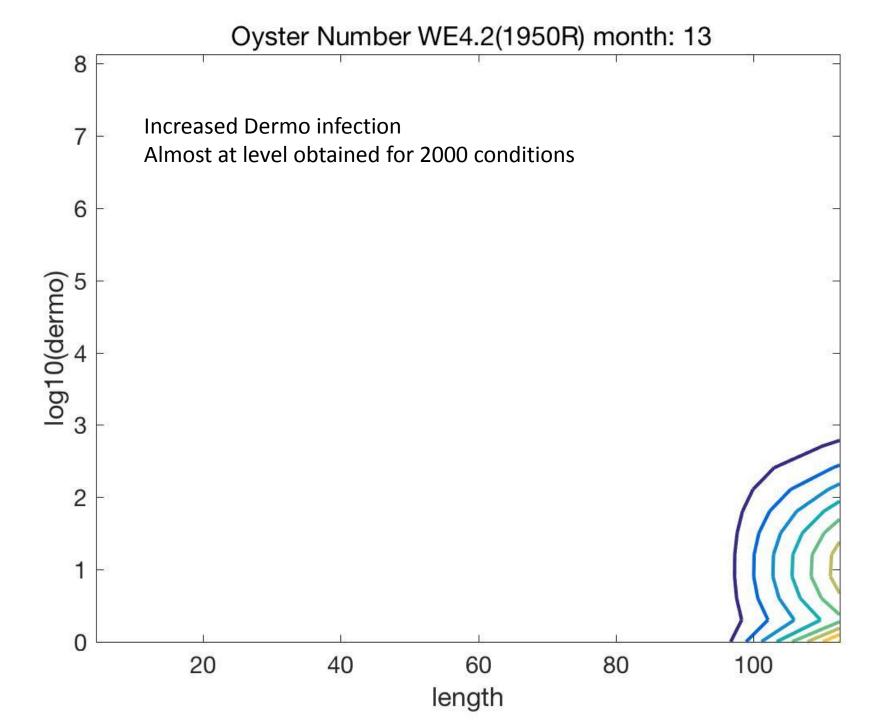
Adjustments

- Used temperature trend to adjust 1900-1905 temperatures to cooler values (3 degree reduction)
- Food values are the sum of POC
- Used 1900 POC values as food input
- Reduced 2000-2005 food by one-third
- More detritus (non-food) in current POC time series



Ovster Number WE4.2(2000) month: 1





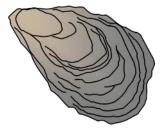
Possible Results

- Increased temperature in 20th century released Dermo pathogen
 - When did temperature exceed pathogen threshold?
- Change in food composition between 1990 and 2000 affects growth rates
 - More 'food' is available but of the wrong form?



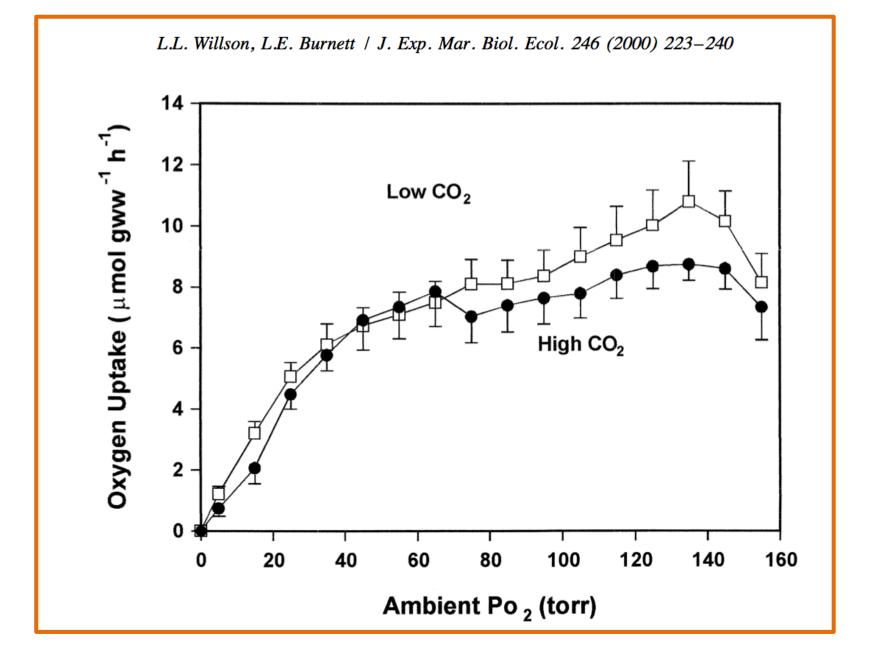
Next Steps?

- Modify oyster model to include explicit oxygen dependencies on physiological processes
- Continue refinement of input time series
- Implement additional simulations to compare past and current conditions and effects
- Input simulated environmental conditions to evaluate effects on Chesapeake Bay oysters



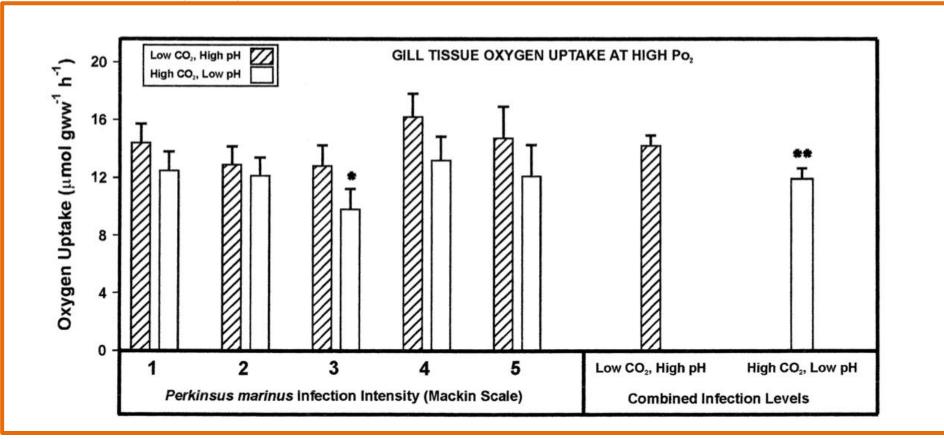
Oxygen

- Post-settlement population respiration, disease intensity, metabolism
- Larvae set survivorship
- Explicit respiration parameterization included for adult oysters – salinity and temperature
- Explicit larval settlement mortality



Hyperbolic relationship that modifies respiration rate

Willson et al. (2000)



No low oxygen effect on disease prevalence; some effect from high CO₂ and low pH

Next Steps?

- Modify oyster model to include explicit oxygen dependencies on physiological processes
- Continue refinement of input time series
- Implement additional simulations to compare past and current conditions and effects
- Input simulated environmental conditions to evaluate effects on Chesapeake Bay oysters

