

Increased Dermo Disease in Chesapeake Bay Oysters Caused by Continued Warming and Nutrient Loading

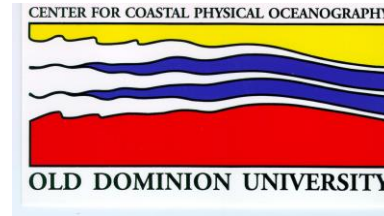
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**CHAMP Project
Meeting
23 June 2021**



Presentation Outline



- Oyster-Dermo model application to Chesapeake Bay
- Simulation results
- Next steps

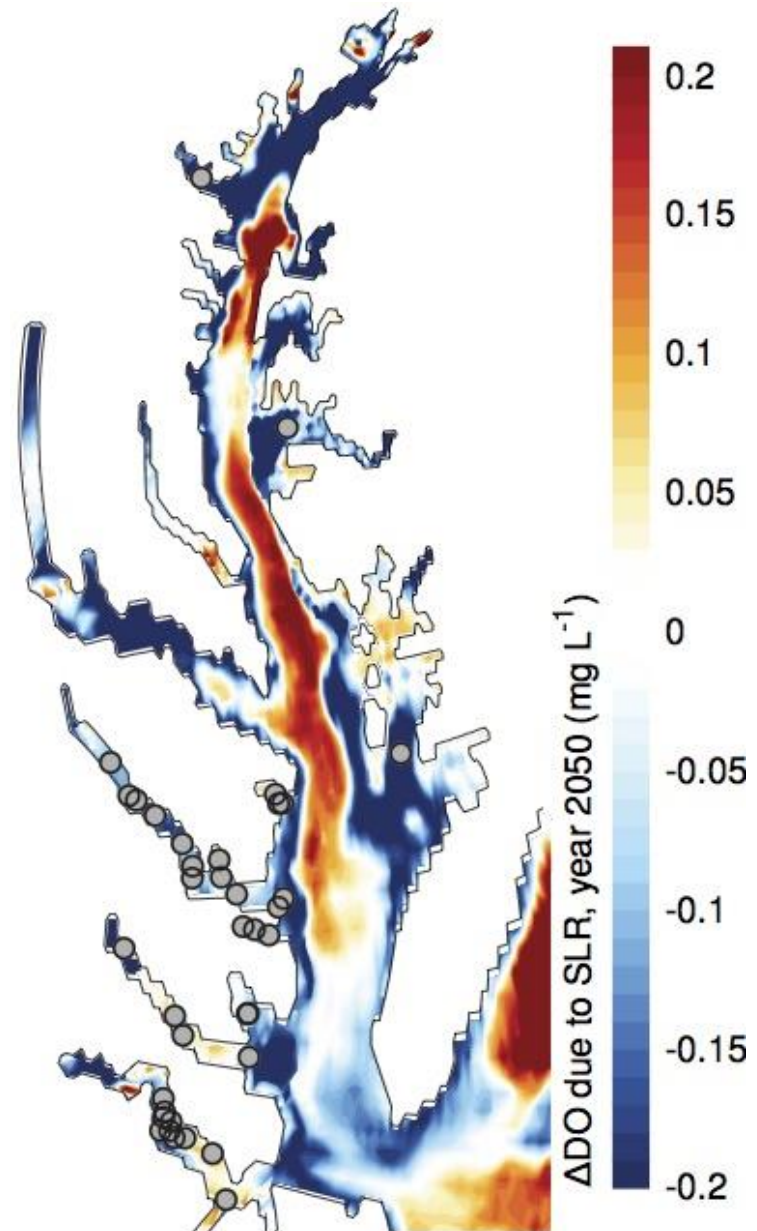
Oyster Model

- Simulate Dermo disease infection intensity and prevalence
- Inputs are temperature, salinity, food from Chesapeake Bay biogeochemical model
 - 1900 -1914
 - 1980 - 2015
- Results presented as Mackin Index; semi-quantitative scale of infection intensity

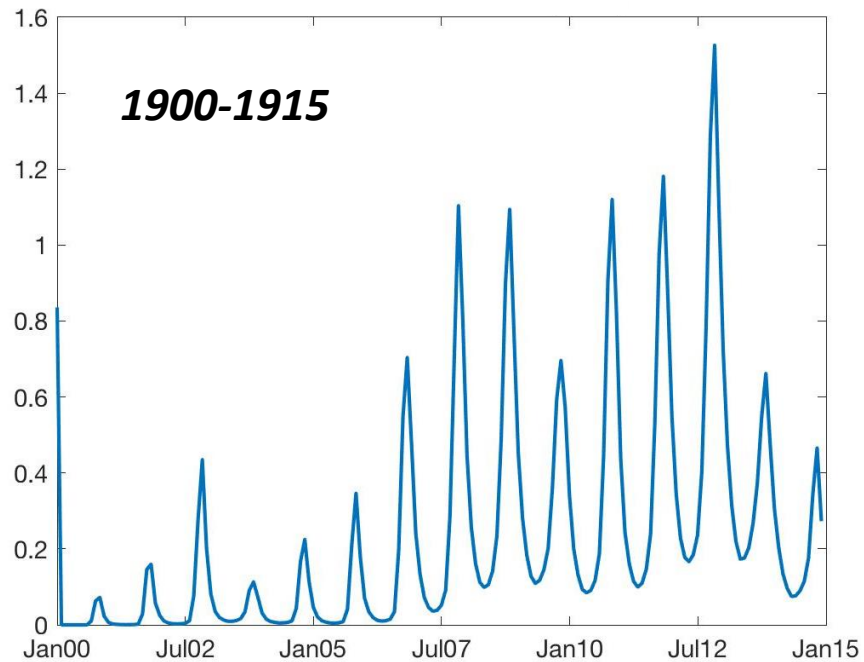


Oyster-dermo model implementation sites

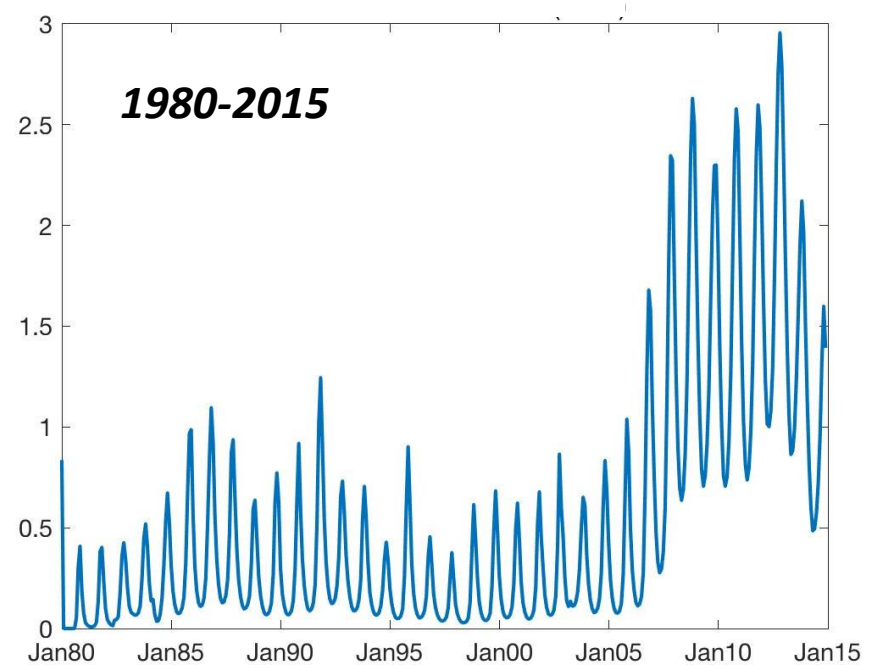
- Coincide with VIMS and Bay program long-term monitoring sites



Mackin Index



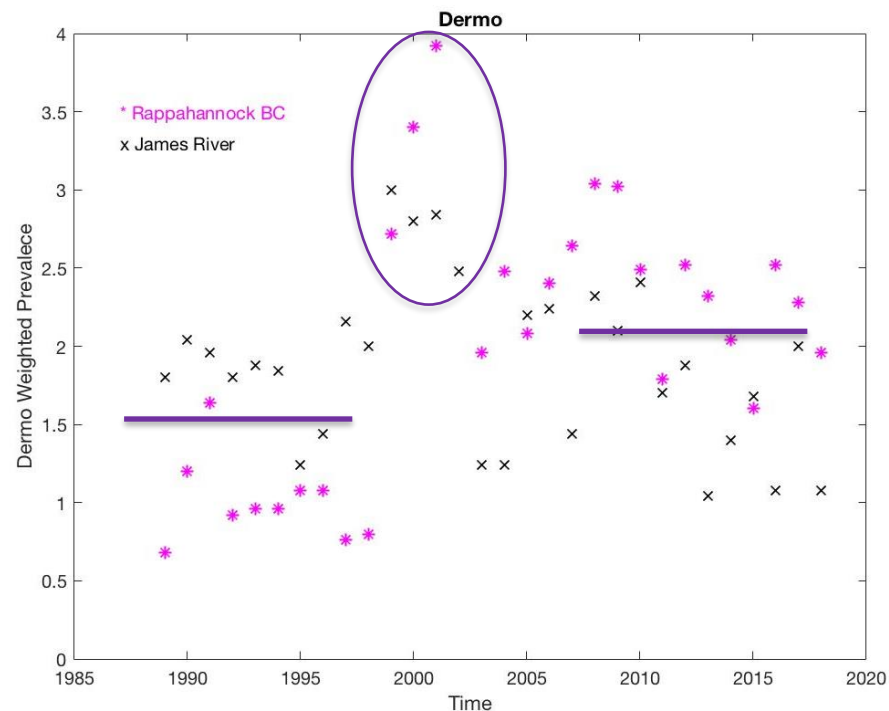
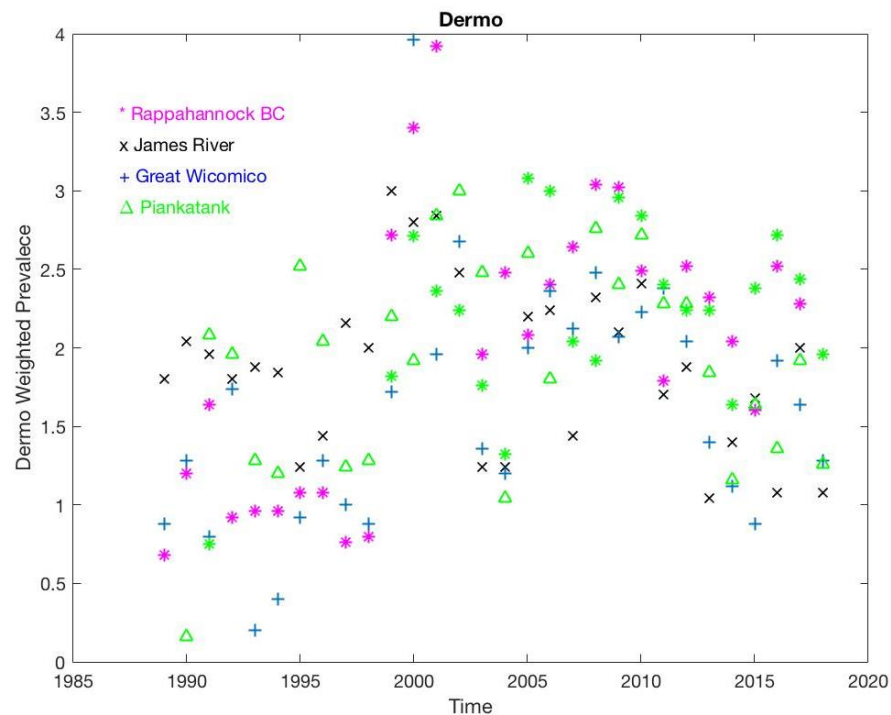
1900-2015

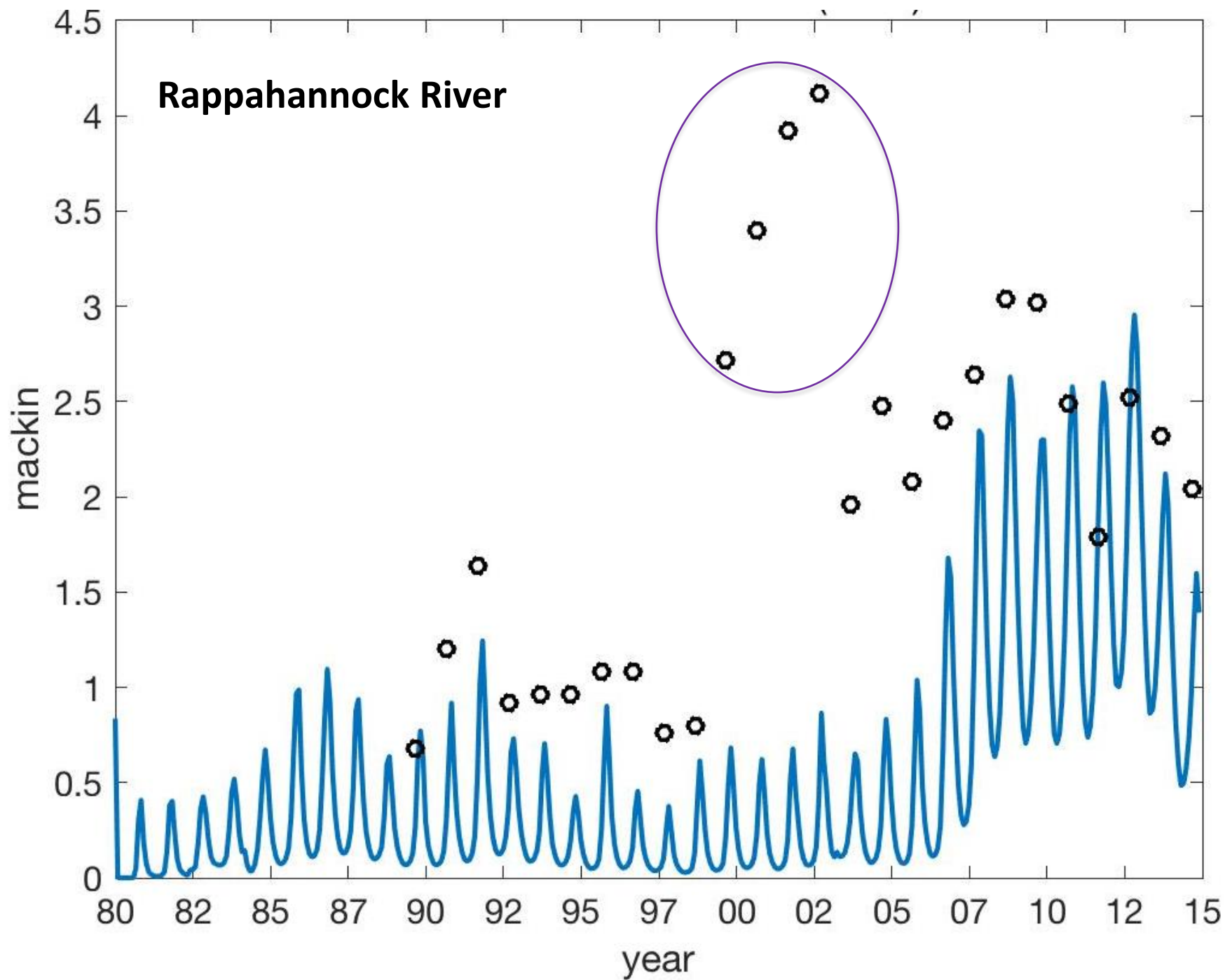


1980-2015

Observed Dermo Weighted Prevalence

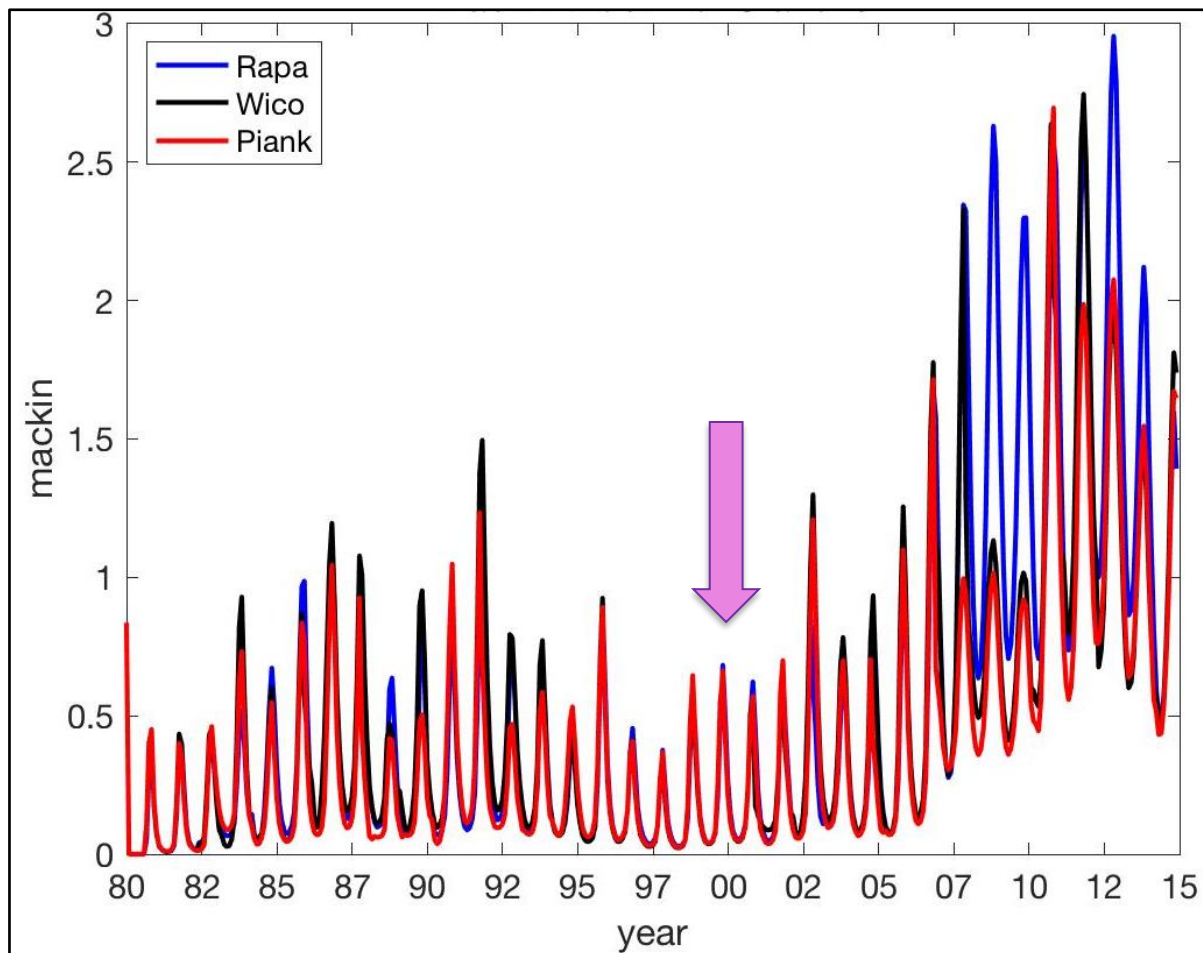
VIMS monitoring data – provided by Ryan Carnegie





Simulated Mackin Index

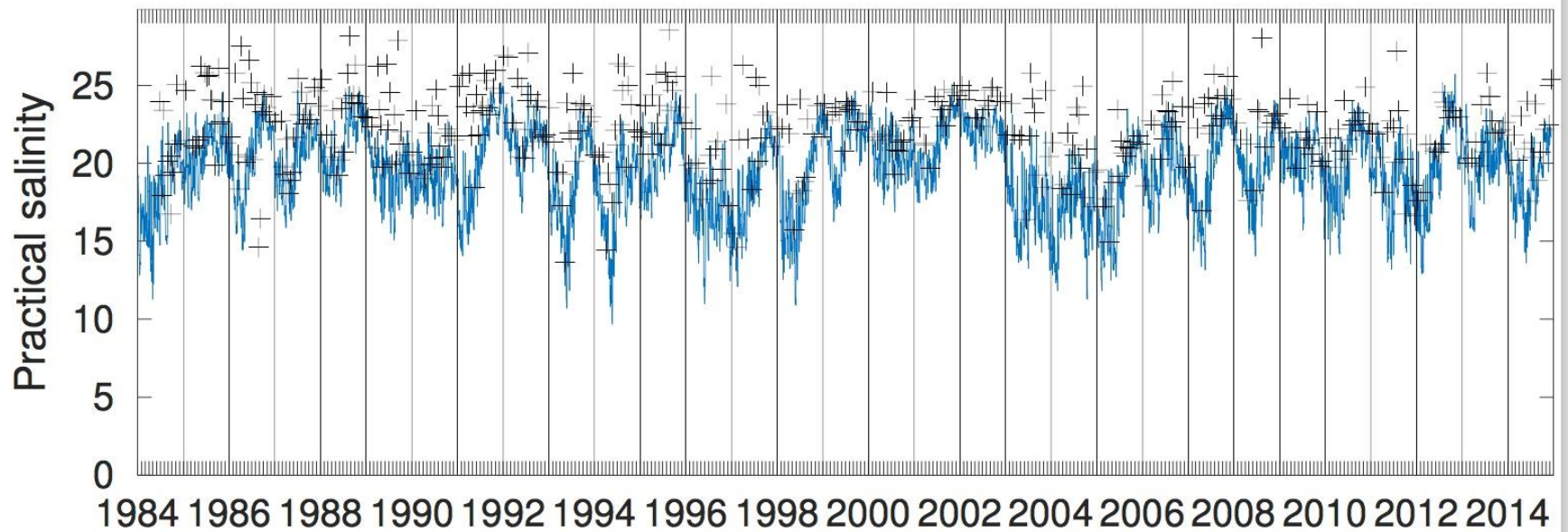
Rappahannock, Wicomico, Piankatank



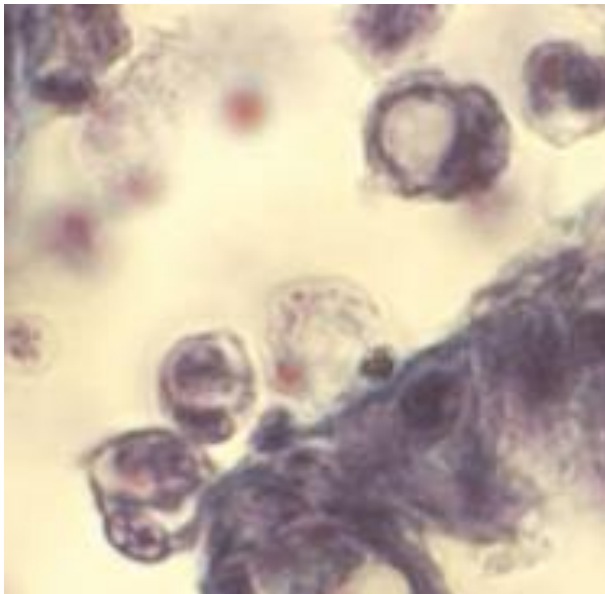
Environmental conditions

- Bottom salinity
- Drought 1999 – 2002

Station 32: WE4.2



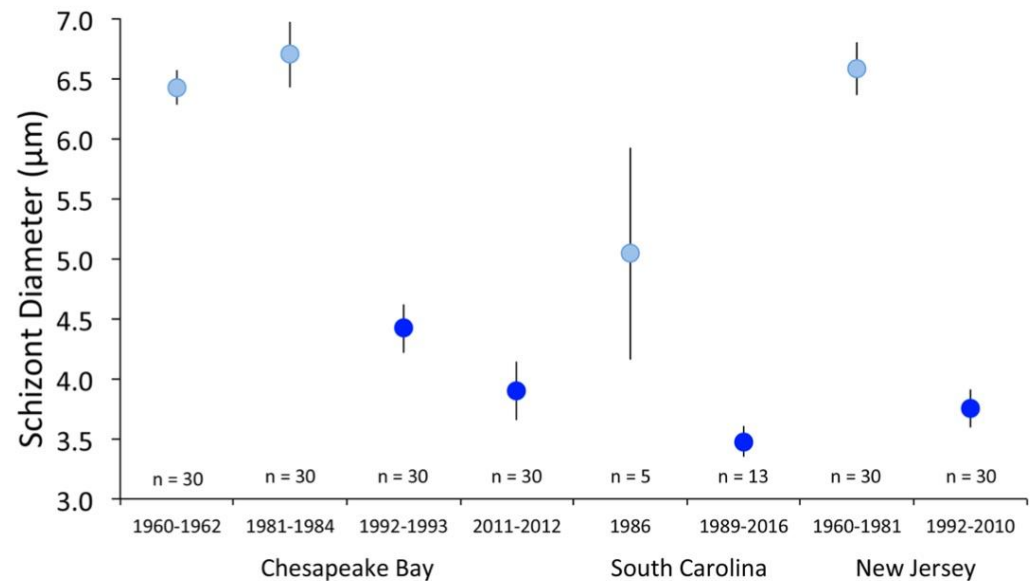
What is happening?



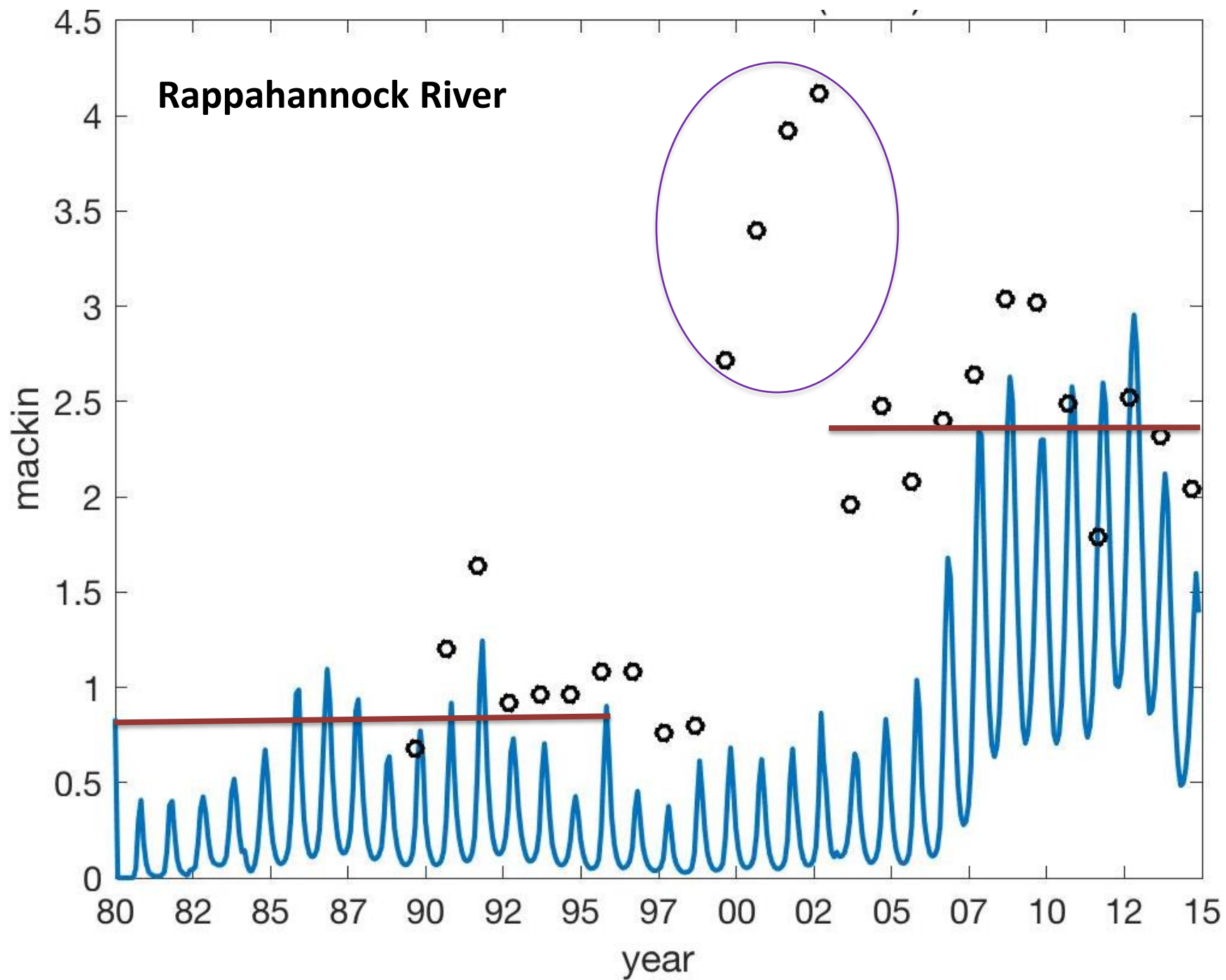
- Environmental conditions may not represent actual conditions
- MSX disease present – dual disease effect
- *“Unprecedented high prevalences and intensities of H. nelsoni were observed in the Great Wicomico, Coan, Yeocomico and Rappahannock Rivers.”* (Calvo & Burreson, 2000)

What is happening?

- Change in dermo pathogen (*Perkinsus marinus*)
- Major phenotypic change producing increased virulence occurred in 1985/1986 epizootic
- Cell diameters smaller by 2011-2012



Carnegie et al. (2021)



Summary

- Temperature higher in 1980s-2000s than early 1900s – released pathogen
- Changed pathogen phenotype and virulence in 1980s
- Occurrence of MSX in 2000 and drought
 - Allowed new pathogen to be established
- Oysters adjusted to a new equilibrium at higher pathogen load
- Rerun simulations
 - modified bottom salinity
 - Increased virulence

