Trends in mycobacteriosis and associated relative mortality in striped bass in Maryland waters of the Chesapeake Bay

Geneviève Nesslage (UMCES CBL)

Jerelle Jesse (UMCES CBL/Univ. of Maine)

Michael Wilberg (UMCES CBL)

Mark Matsche (Maryland DNR)

Howard Townsend (NOAA)

Chunqi Shen (UMCES CBL/SUST)

Jeremy Testa (UMCES CBL)

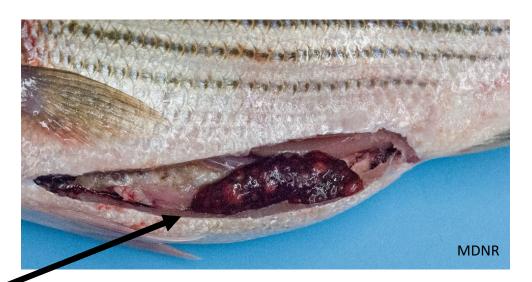






Mycobacteriosis

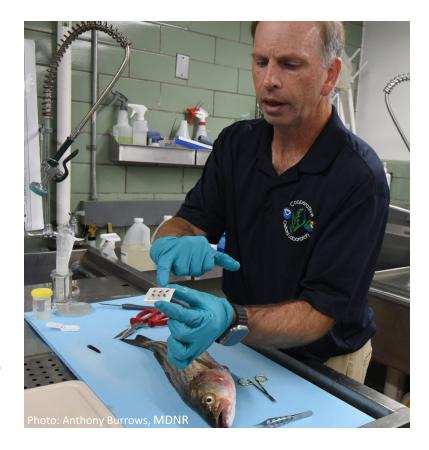
- Chronic disease from bacteria of the genus Mycobacterium
- Identified in several fish species
- Symptoms include
 - Nodules on internal organs
 - Scale loss, pigment
 changes, external
 lesions





Mycobacteriosis in Chesapeake Bay striped bass

- Disease first described in 1997, archived samples as far back as 1984 have tested positive (Jacobs et al. 2009)
- Disease prevalence high in Chesapeake Bay, low in coastal ocean (Matsche et al. 2010)



Mycobacteriosis affects striped bass population dynamics

- Decreased growth and earlier/smaller size at maturity (Latour et al. 2010, Gervasi et al. 2019)
- Relative survival of diseased fish lower (Gauthier et al. 2008, Groner et al. 2018)
- Relative survival of severely diseased fish half that of healthy fish (Hoenig 2017)



Maryland Department of Natural Resources Fish Health Program

1998 - present Opportunistic and directed sampling with various gears



Sex, weight, condition, length/age



Disease presence and severity

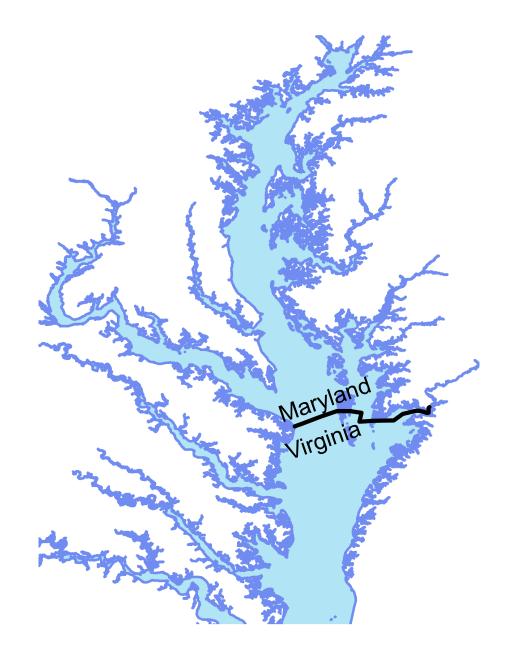






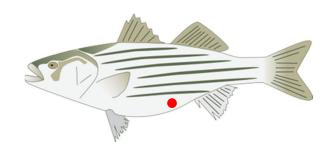
Objectives

 Quantify trends in mycobacteriosis in striped bass in Maryland waters of the Chesapeake Bay

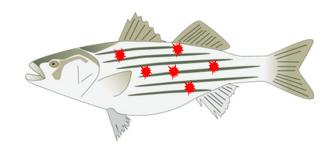


Objectives

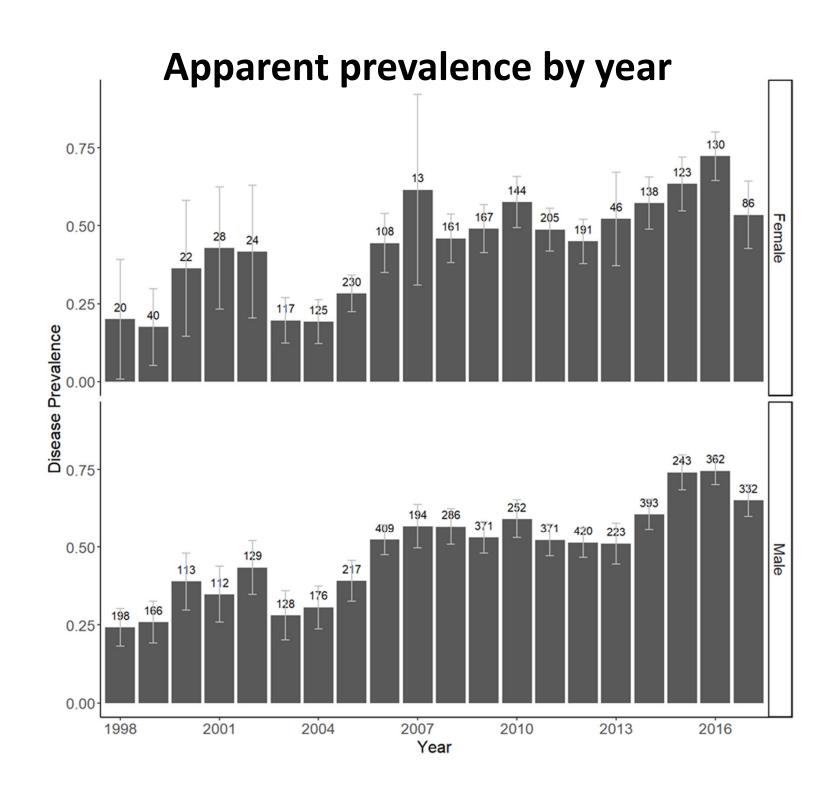
2. Model relationship between disease apparent prevalence and potential disease drivers



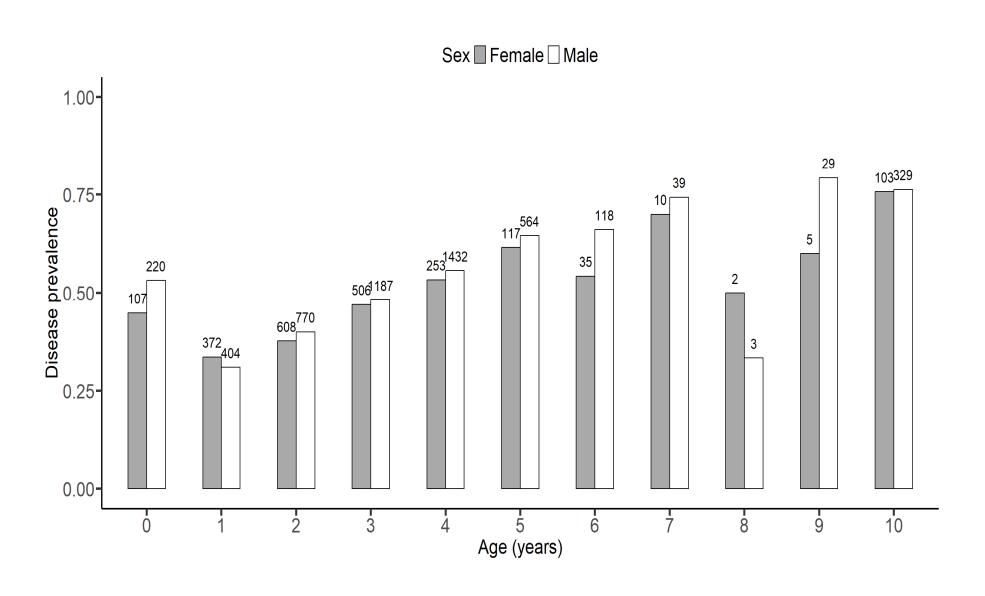
Model relationship between disease severity and potential disease drivers



Objective 1 : Quantify trends



Apparent prevalence by age



Objectives 2-3: Identify potential drivers of disease

Potential drivers

• Fish condition: if fish in poor condition due to malnutrition, fish more susceptible to disease

• Environmental: if water conditions (e.g., temperature & dissolved oxygen) are poor, fish may be stressed or change feeding behavior

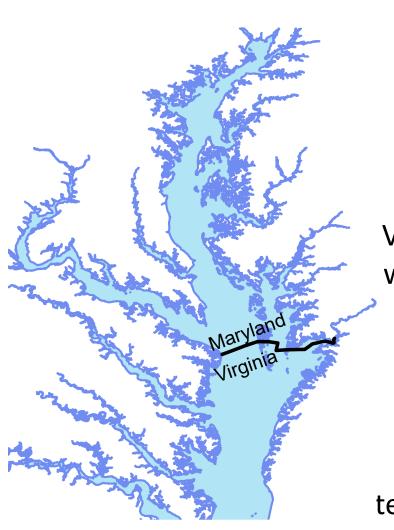
Potential drivers

• Fish condition: if fish in poor condition due to malnutrition, fish more susceptible to disease

Fulton's K – condition metric (total weight/length³)

• Environmental: if water conditions (e.g., temperature & dissolved oxygen) are poor, fish may be stressed or change feeding behavior

Coupled hydrodynamic-biogeochemical model (ROMS-RCA) – 1998-2015



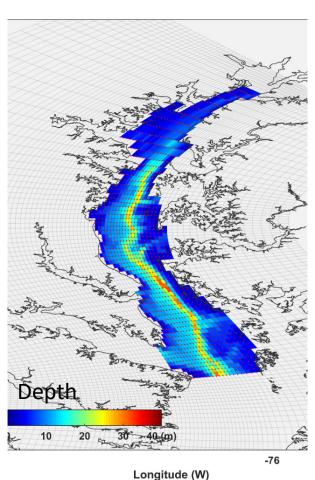
Annual summer average:

Hypoxic Volume:

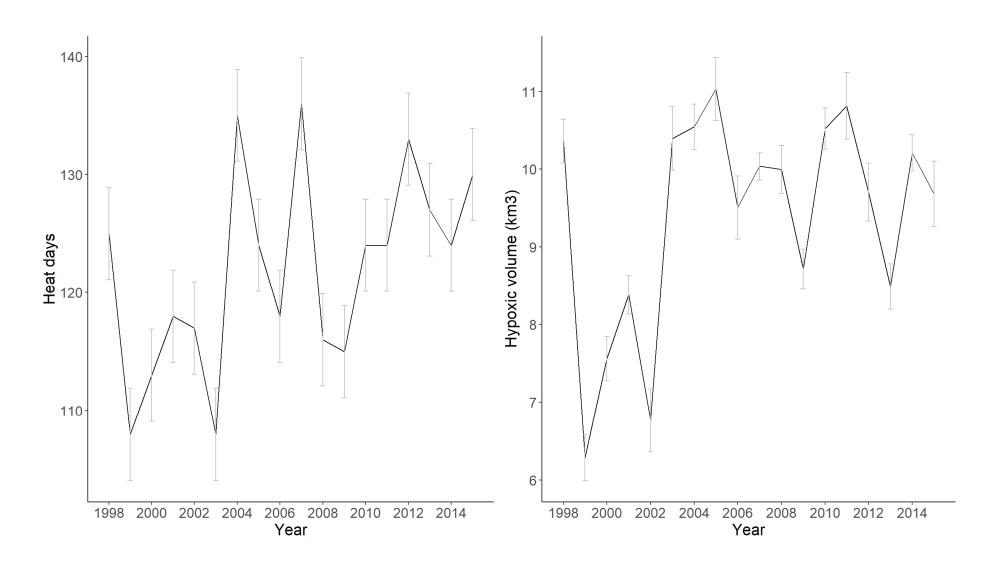
Volume of bottom water with DO < 3 mg/L

Heat Days:

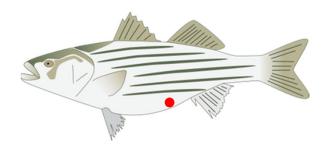
Number of days surface water temperature > 25 C



ROMS-RCA estimates



Apparent prevalence model ?

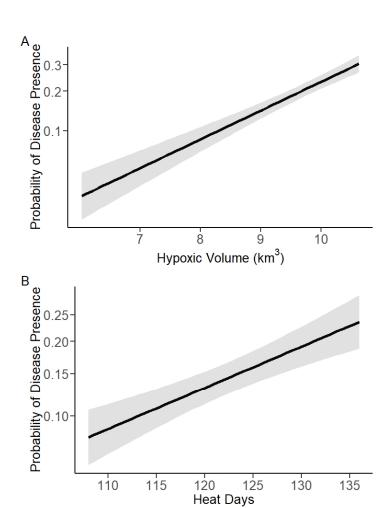


Binomial GLM

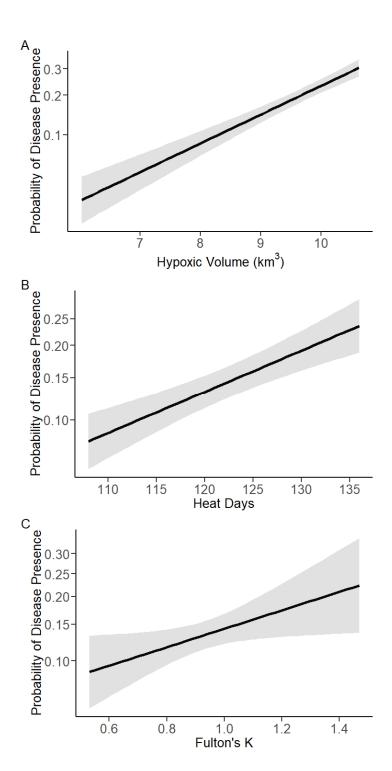
- -Diseased (+ mycobacteria in spleen samples) or not
- -age 0-1 pre-migratory fish (known residents)
- -1998-2015

 $\hat{Y} = \beta_0 + \beta_1 Hypoxic Volume + \beta_2 Heat Days + \beta_3 Fulton's K$

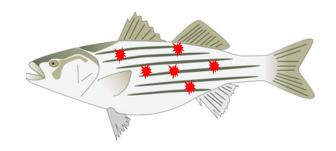
 Disease presence in premigratory fish increased with increasing hypoxic volume and heat days



- Disease presence in premigratory fish increased with increasing hypoxic volume and heat days
- Disease presence for pre-migratory fish increased with increasing fish condition



Disease severity model

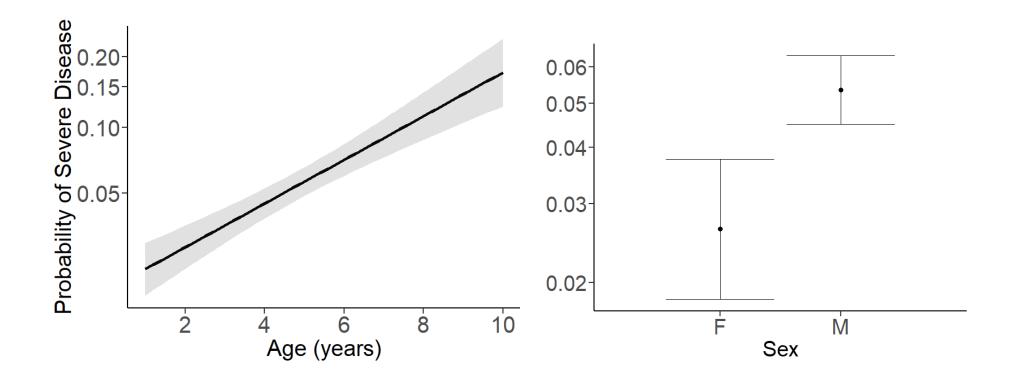


Binomial GLM

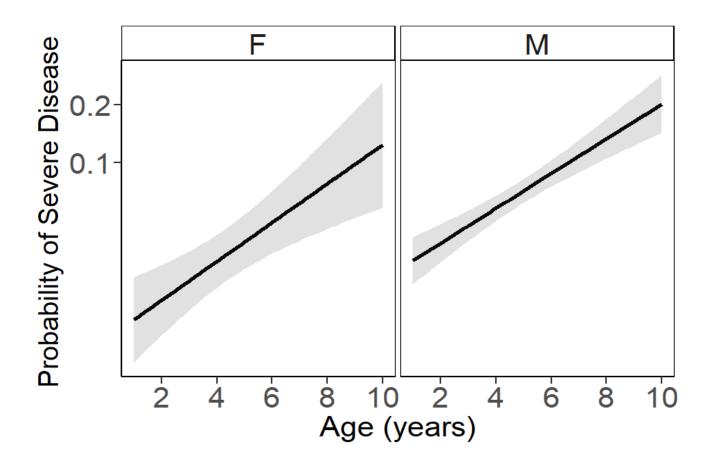
- -Severe external lesion severity score (yes/no)
- -age 1+ fish
- -2007-2017

$$\hat{Y} = \beta_0 + \beta_1 Hypoxic Volume + \beta_2 Heat Days + \beta_3 Fulton's K + \beta_4 Age + \beta_5 Sex + \beta_6 Age * Sex,$$

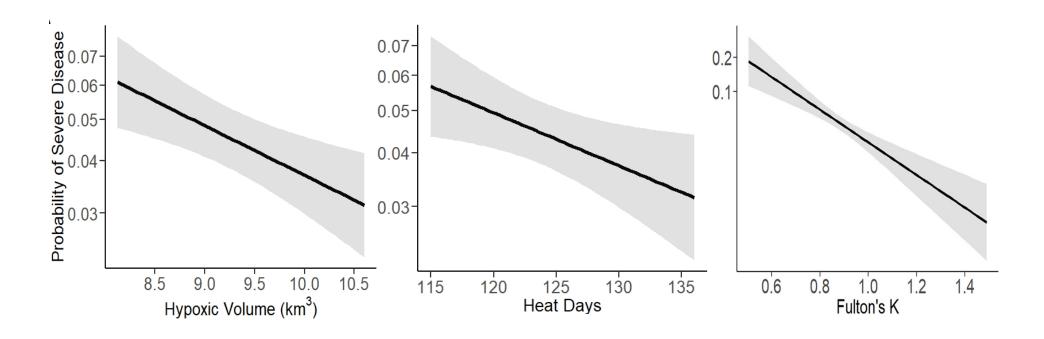
$$Probability = \frac{e^{\beta_0 + \beta_1 Hypoxic Volume + \beta_2 Heat \ Days + \beta_3 Fulton's \ K + \beta_4 Age + \beta_5 Sex + \beta_6 Age * Sex}}{1 + e^{\beta_0 + \beta_1 Hypoxic Volume + \beta_2 Heat \ Days + \beta_3 Fulton's \ K + \beta_4 Age + \beta_5 Sex + \beta_6 Age * Sex}}$$



- Occurrence of severe disease increased with age
- Males more likely to be severely diseased, but...

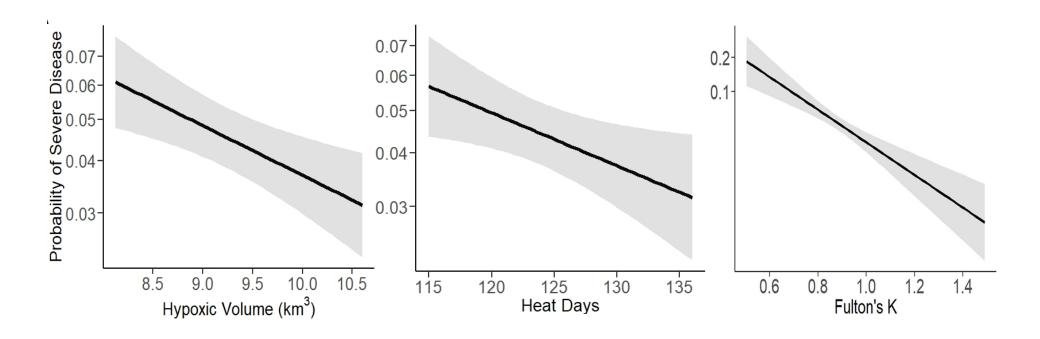


Similar progression of disease with age for both sexes

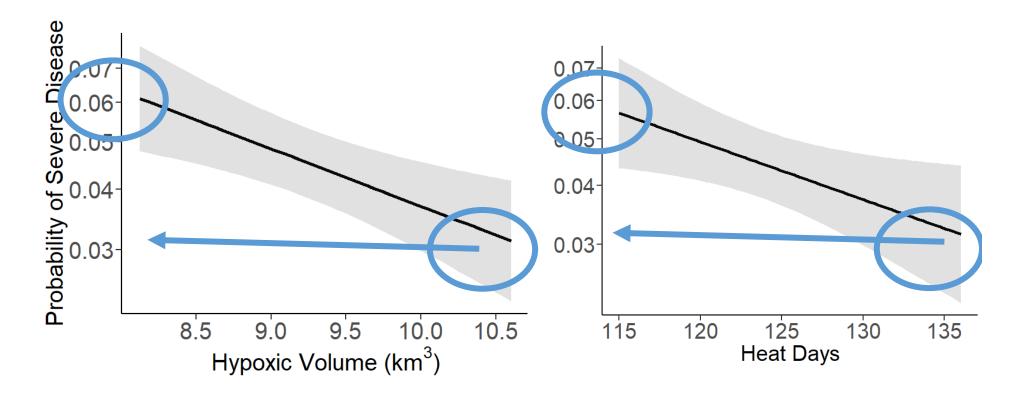


Counterintuitive results:

- Occurrence of severe disease decreased with increasing hypoxic volume and heat days
- Occurrence of severe disease decreased with increasing fish condition



- Evidence of mortality among severely diseased fish?
- If dead, can't be caught in survey!



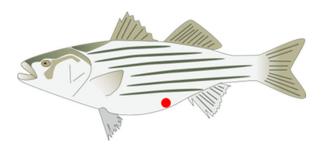
- If severely diseased fish are dying at a higher rate, we can estimate their mortality relative to other fish
- Implies approx. doubling of natural mortality for severely diseased fish
- Similar to other published studies (Gauthier et al. 2008, Hoenig et al. 2017, Groner et al. 2018)

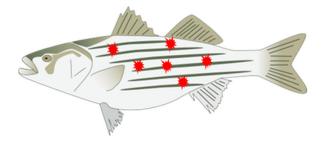
Deeper dive into fish condition

- Modeled prevalence using only fish with no disease vs mild disease, and found fish condition (Fulton's K) no longer a significant covariate but hypoxic volume and heat days remained significant covariates
- Modeled fish condition as a function of severity (all ages), but disease severity alone could not predict fish condition
- Poor condition may contribute to disease contraction and progression in some fish
- However, Maryland data suggest fish condition is more likely a symptom of disease than a cause

Conclusions

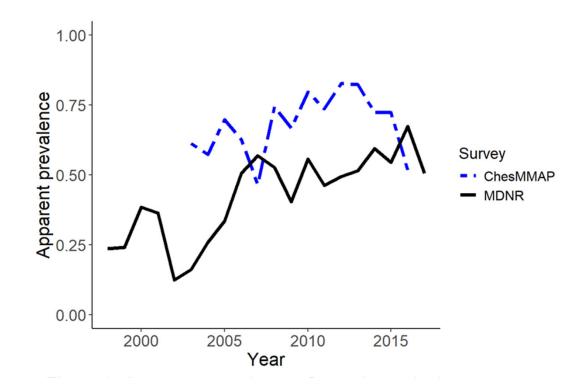
- Mycobacteriosis is widespread in the Maryland portion of the Chesapeake Bay
- Disease prevalence and severity in Maryland striped bass is related to poor water quality and fish condition
- Relative mortality for severely diseased fish implied by our models <u>doubled</u> across the range of low oxygen and water temperature conditions examined in this study





Next steps

- Team formed: CBL, VIMS, Maryland DNR, NOAA, URI/RIDEM, ASMFC
- Pursuing funding to:
 - Comprehensively examine all myco data across Chesapeake Bay
 - Develop age-specific mycobacteriosis trends
 - Estimate age-specific trends in natural mortality (Schonfeld 2023)



Jesse et al. 2024 (in proofs). Quantifying trends in and potential drivers of mycobacteriosis in Atlantic Striped Bass in Maryland waters of the Chesapeake Bay.

Transaction of the American Fisheries Society.

Genny Nesslage nesslage@umces.edu

