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# The Snow Crab IBM & BCM: Basis for Potential ESP Indicators?

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May, 2022

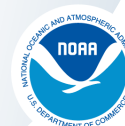
# Acknowledgments

- Contributors

- Michael Torre, Christine Stawitz, Bob Foy, Cody Szuwalski
- Al Hermann, Kelly Kearney, Wei Cheng
- Erin Fedewa, Louise Copeman, Jon Richar
- Kirstin Holsman, Kerim Aydin, Anne Hollowed

- Funding:

- Magnuson-Stevens Act
- FATE



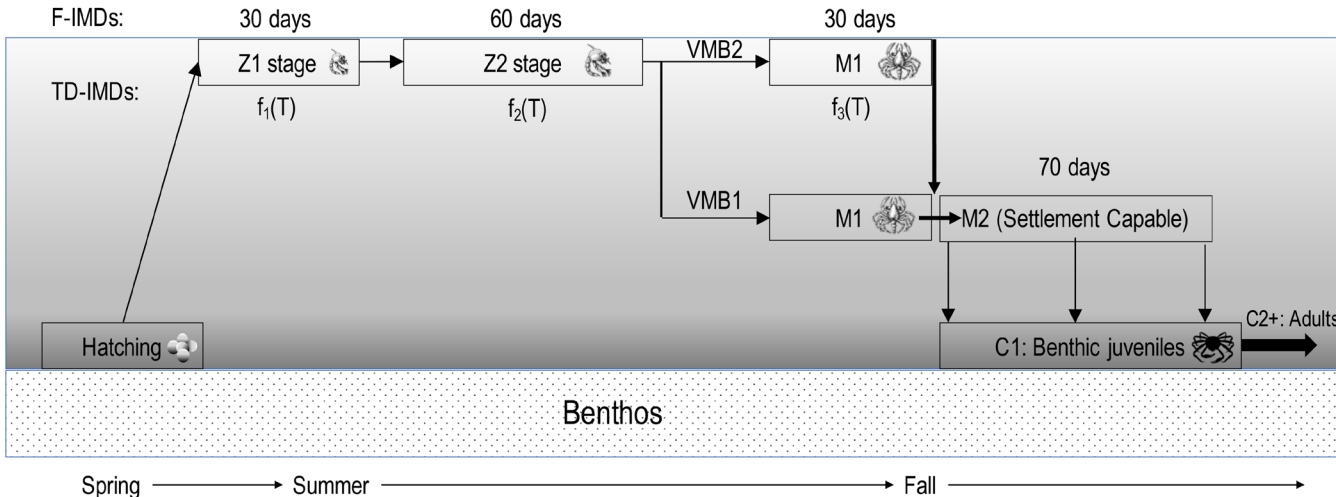
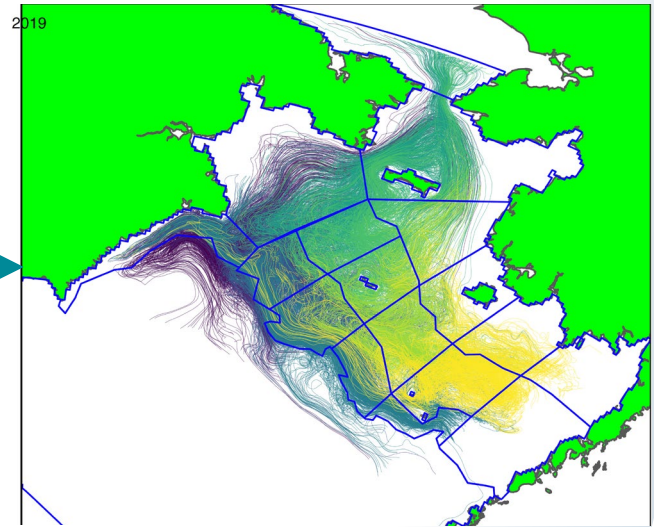
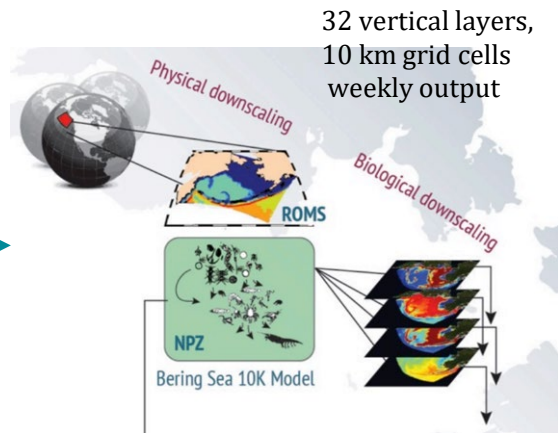
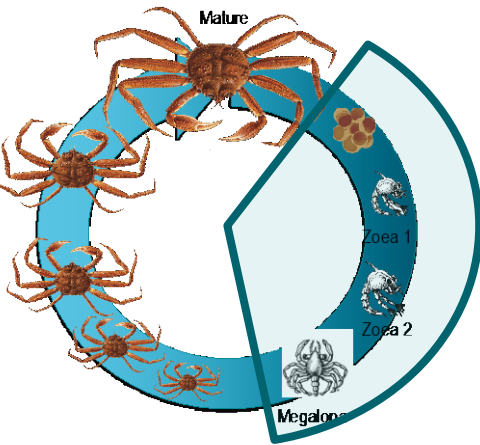
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# What am I talking about?

- IBM: individual-based model
  - tracks simulated individuals through larval, postlarval life stages from hatch location to settlement
  - movement patterns
    - ROMS ocean currents
    - vertical movement behavior
  - development (growth)
    - temperature dependence of molt duration (lab experiments)
    - in situ ROMS water column temperatures
  - mortality: assumed rates
  - settlement in benthic nursery habitats
    - (assumed) depth range + temperature range
- BCM: benthic cohort model (follow-on to IBM)
  - tracks benthic instar abundance in ROMS grid by settlement cohort
  - in situ ROMS bottom temperatures
    - temperature dependence of molt duration
    - temperature dependent mortality rates (think “cod predation”)



# Snow crab IBM

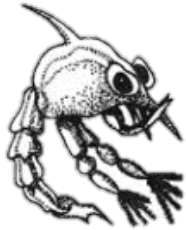


# Pelagic life stages

Z1



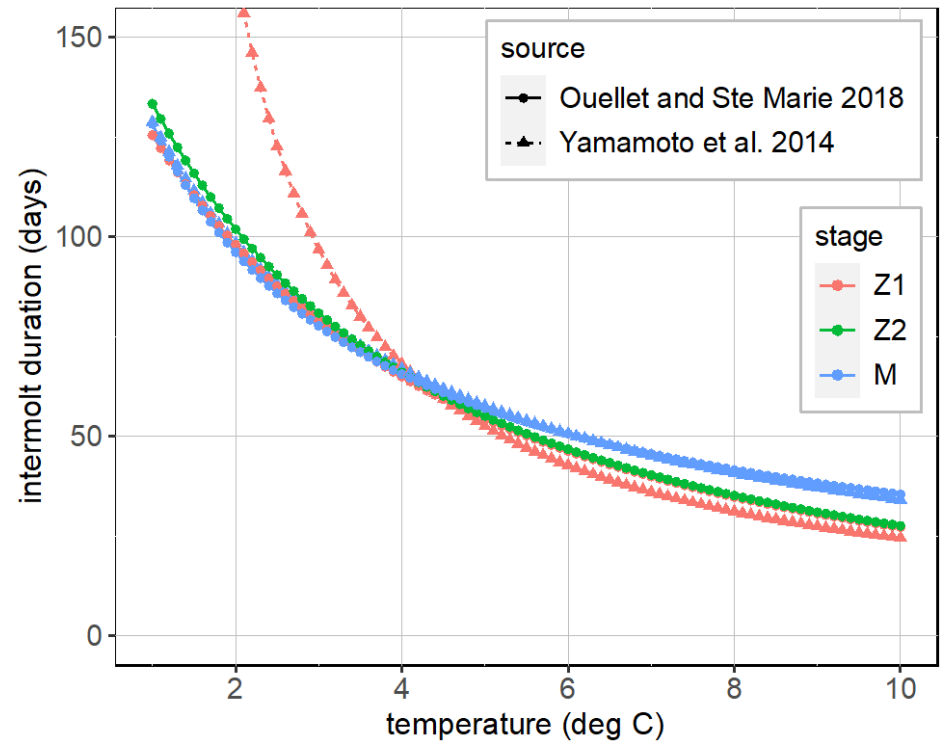
Z2



M



- temperature-dependent or fixed intermolt durations (IMDs)
- “swim” up or down into preferred depth ranges
  - zoea: 0 – 20 m (Incze 1987, Ouellet & Ste Marie 2018)
  - megalopae: 50 – 150 m (Lovrich 1995, Ouellet & Ste Marie 2018)
  - undergo vertical random walks within preferred depth ranges
  - no active horizontal movement (just drift)
- settlement habitat (suitable benthic nursery habitat)
  - assumed depth ranges (25-200 m)
  - assumed temperatureranges



# Model start

Annual model runs

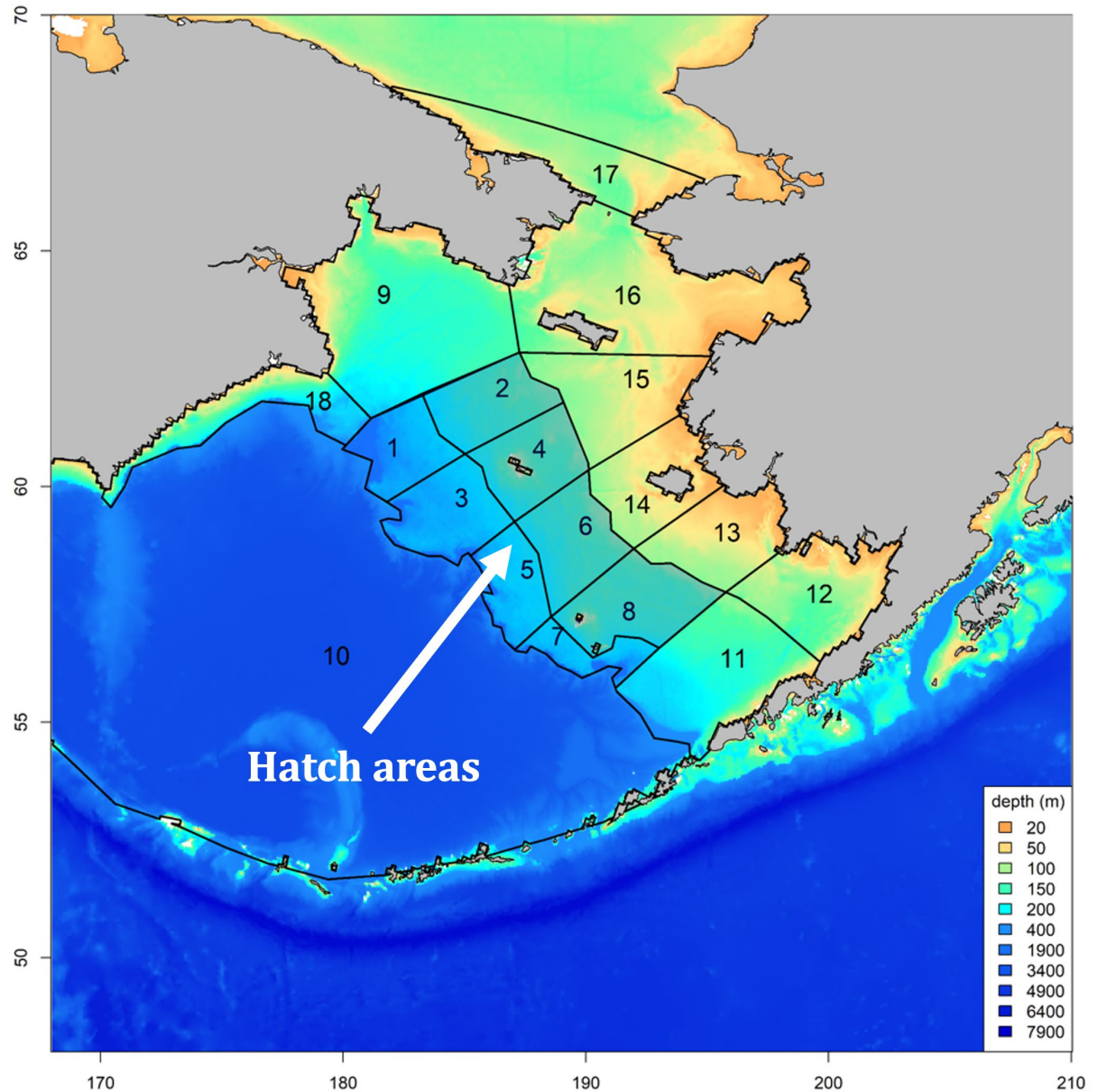
- 1971-2019
- start: hatch date
- end: Dec. 1

5 hatch dates

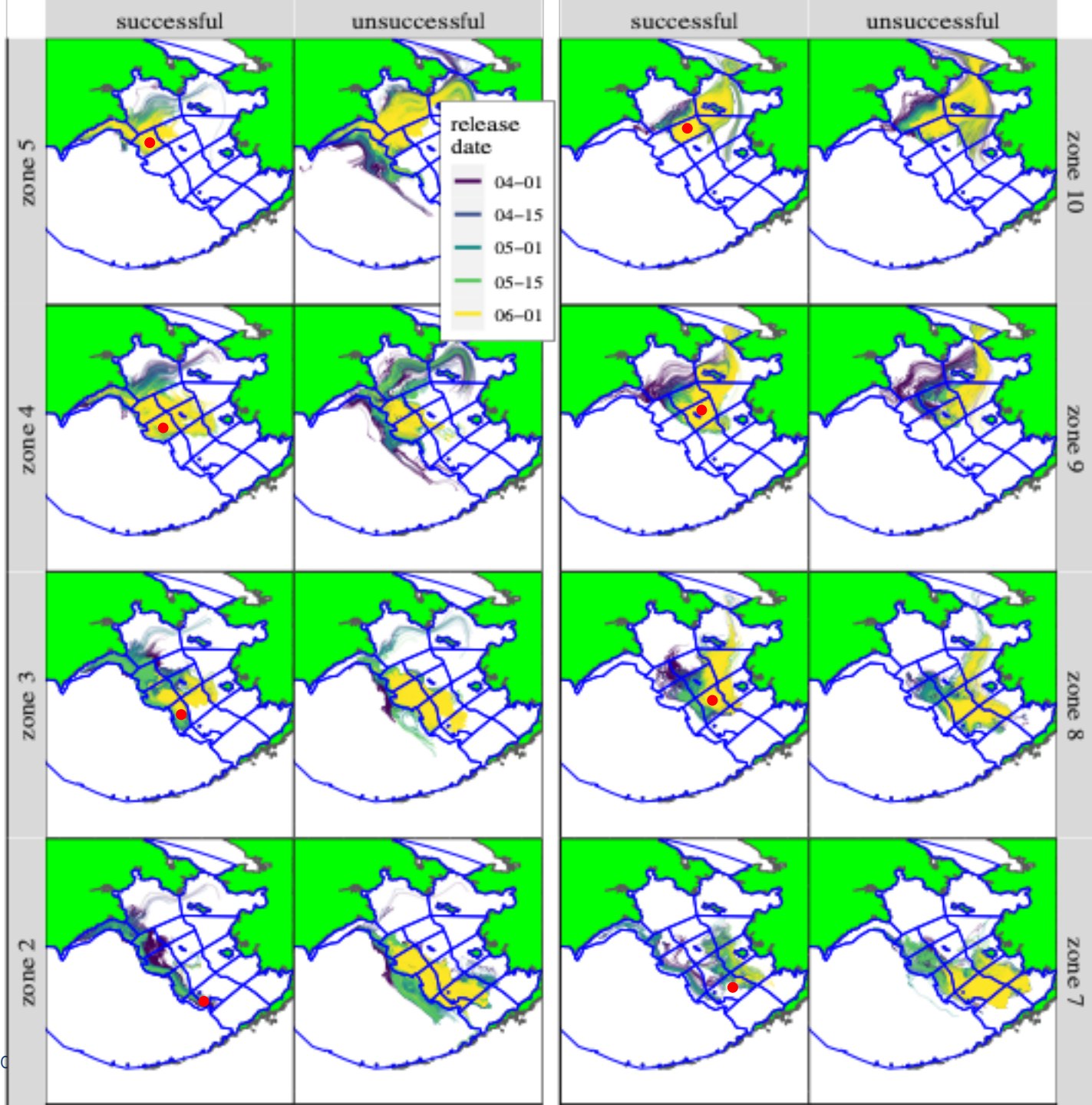
- Apr 1, 15
- May 1, 15
- Jun 1

32,651 individuals/hatch

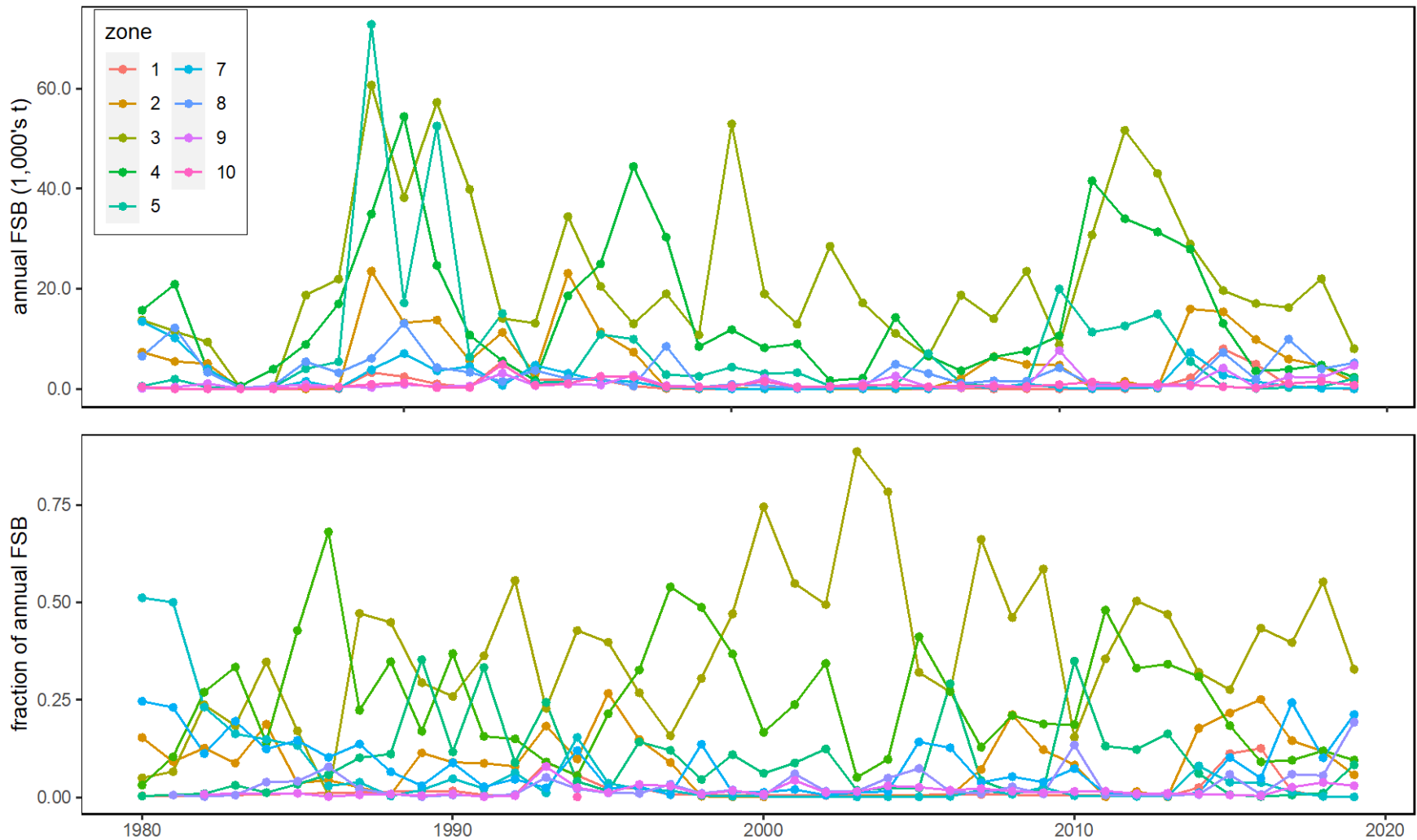
- 3 km spacing



# Example Trajectories

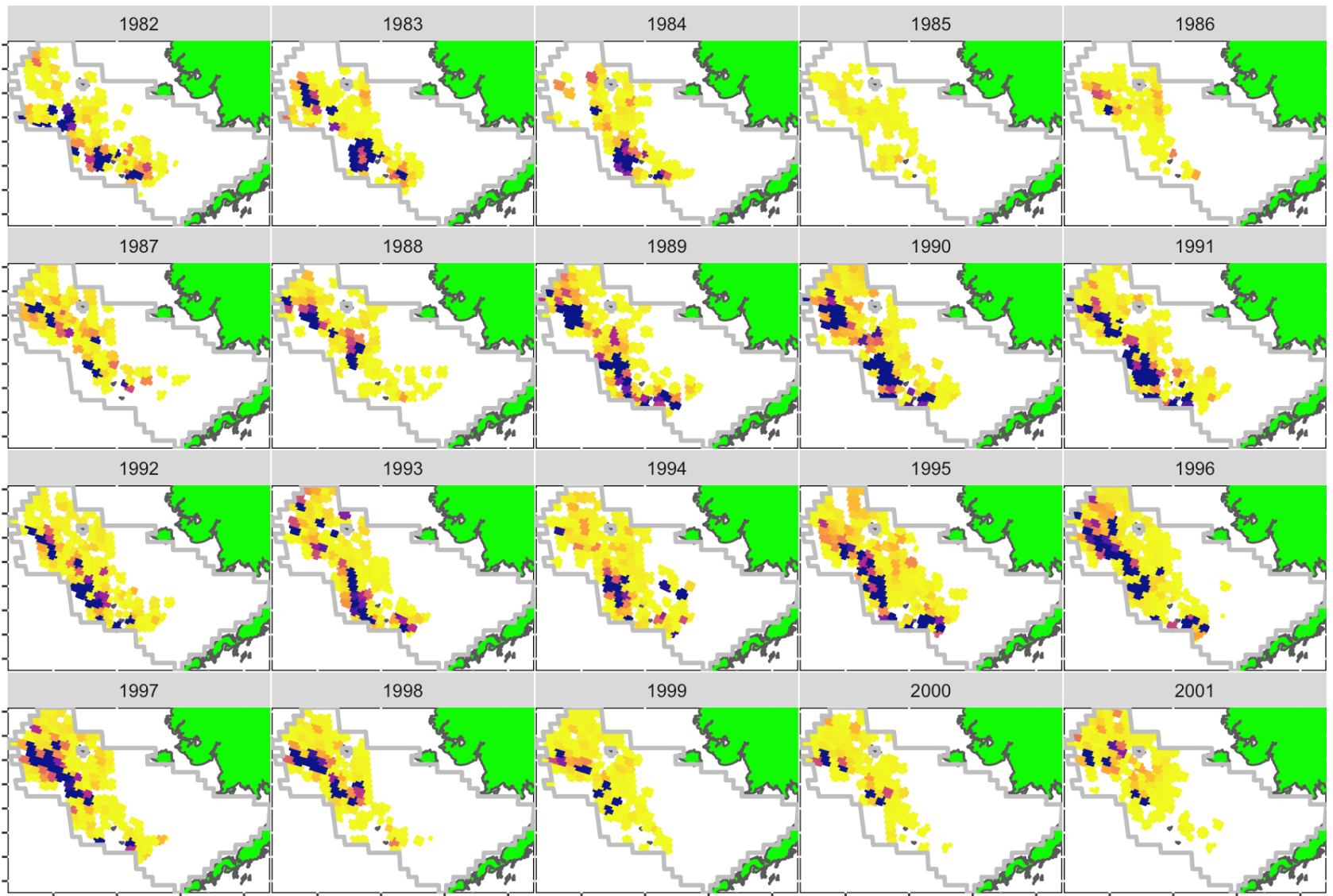


# Adding in variability in egg production: spatiotemporal variation old shell female biomass





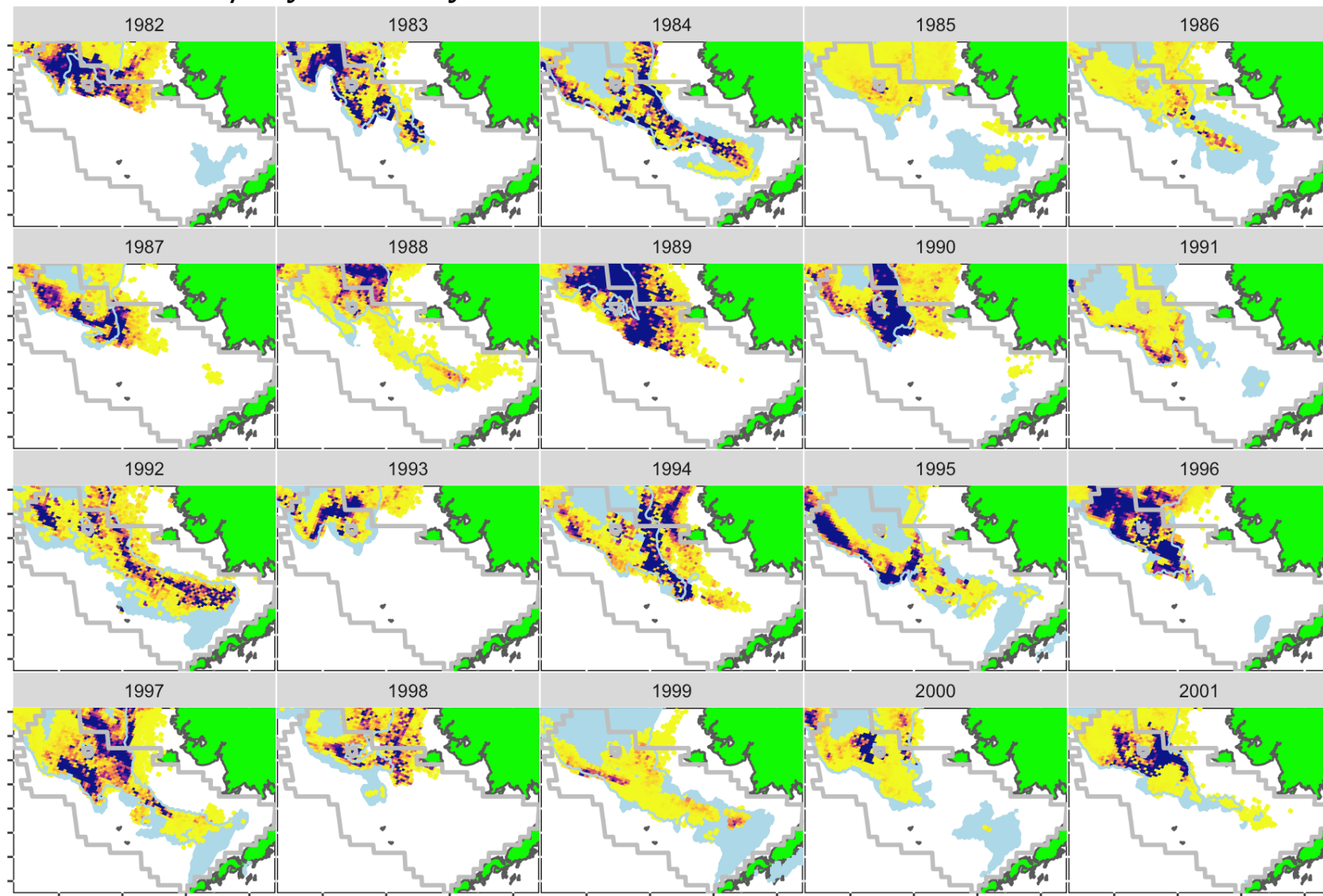
# Hatch Patterns: 1982-2001



# Settlement Patterns: 1982-2001

assumed: 0%/day mortality

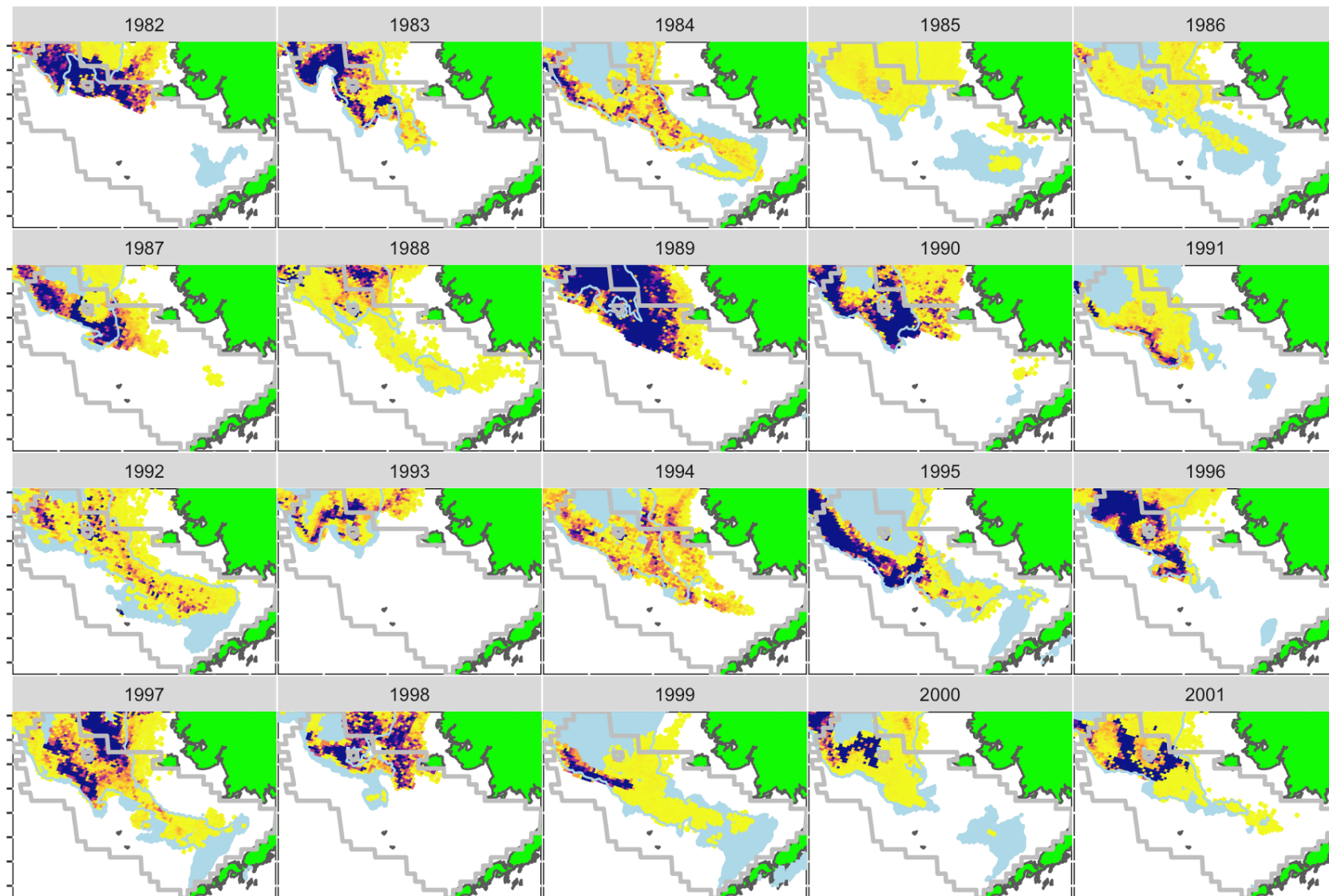
Base Model



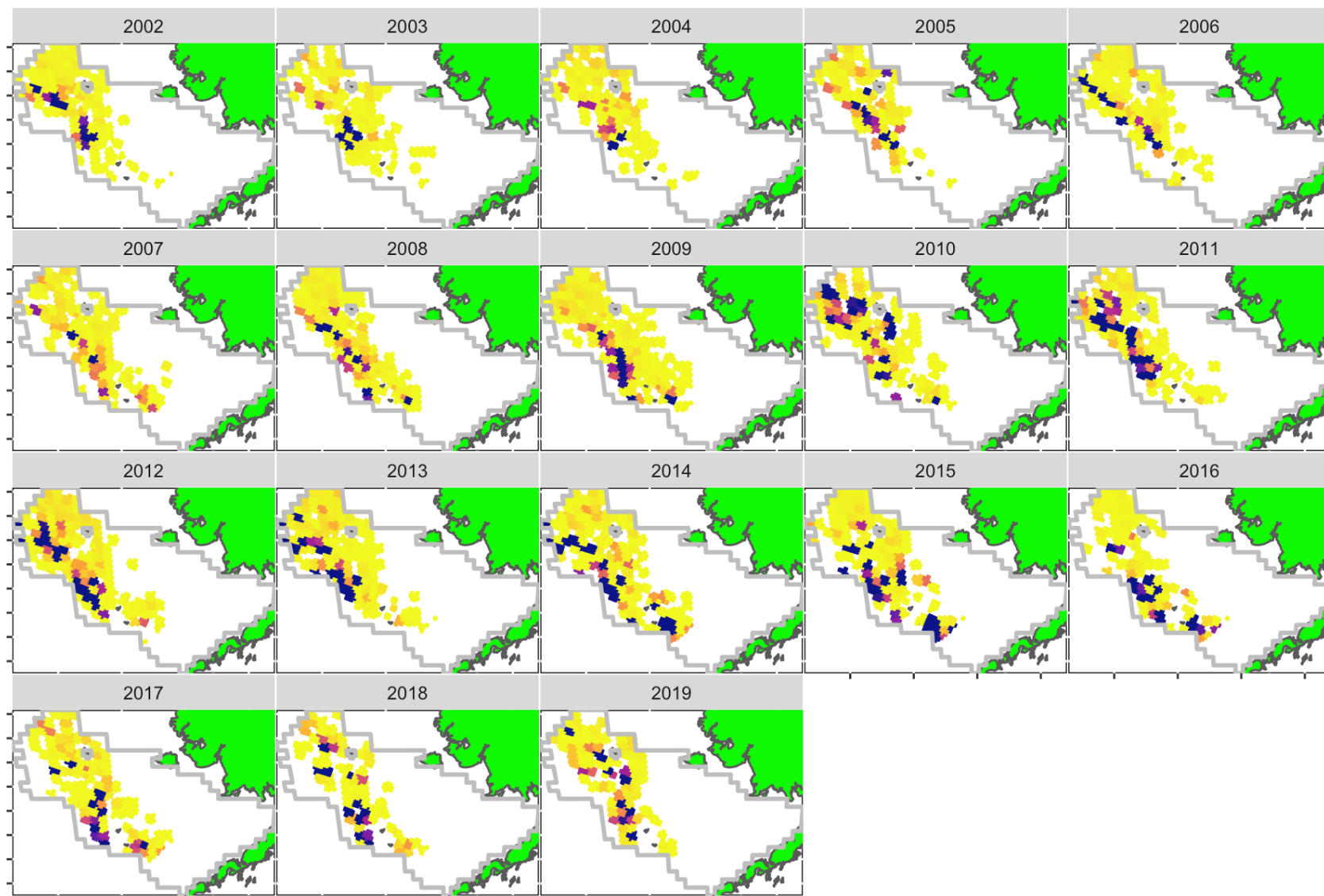
# Settlement Patterns: 1982-2001

assumed: 8%/day mortality

Base Model



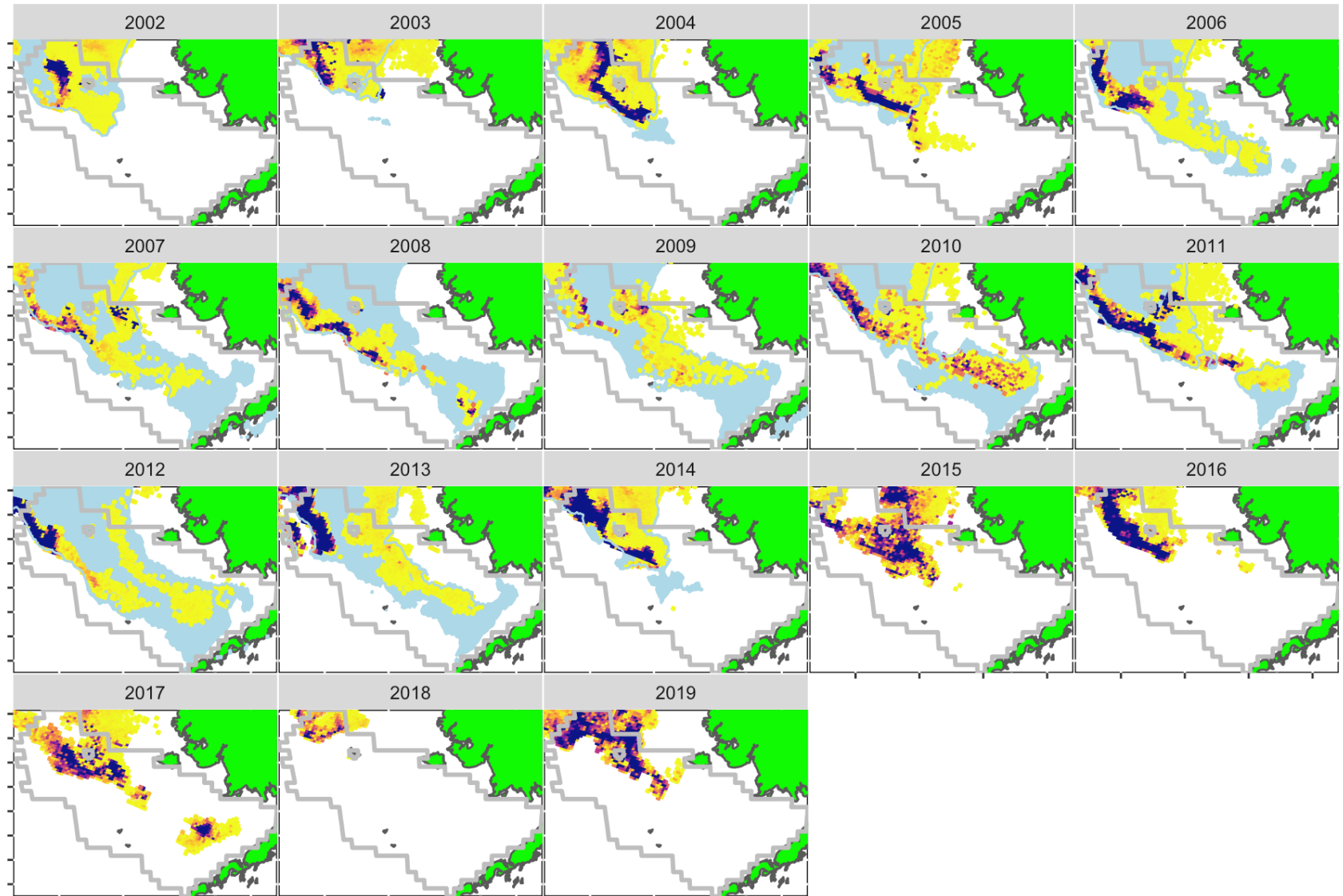
# Hatch Patterns: 2002-2019



# Settlement Patterns: 2002-2019

assumed: 0%/day mortality

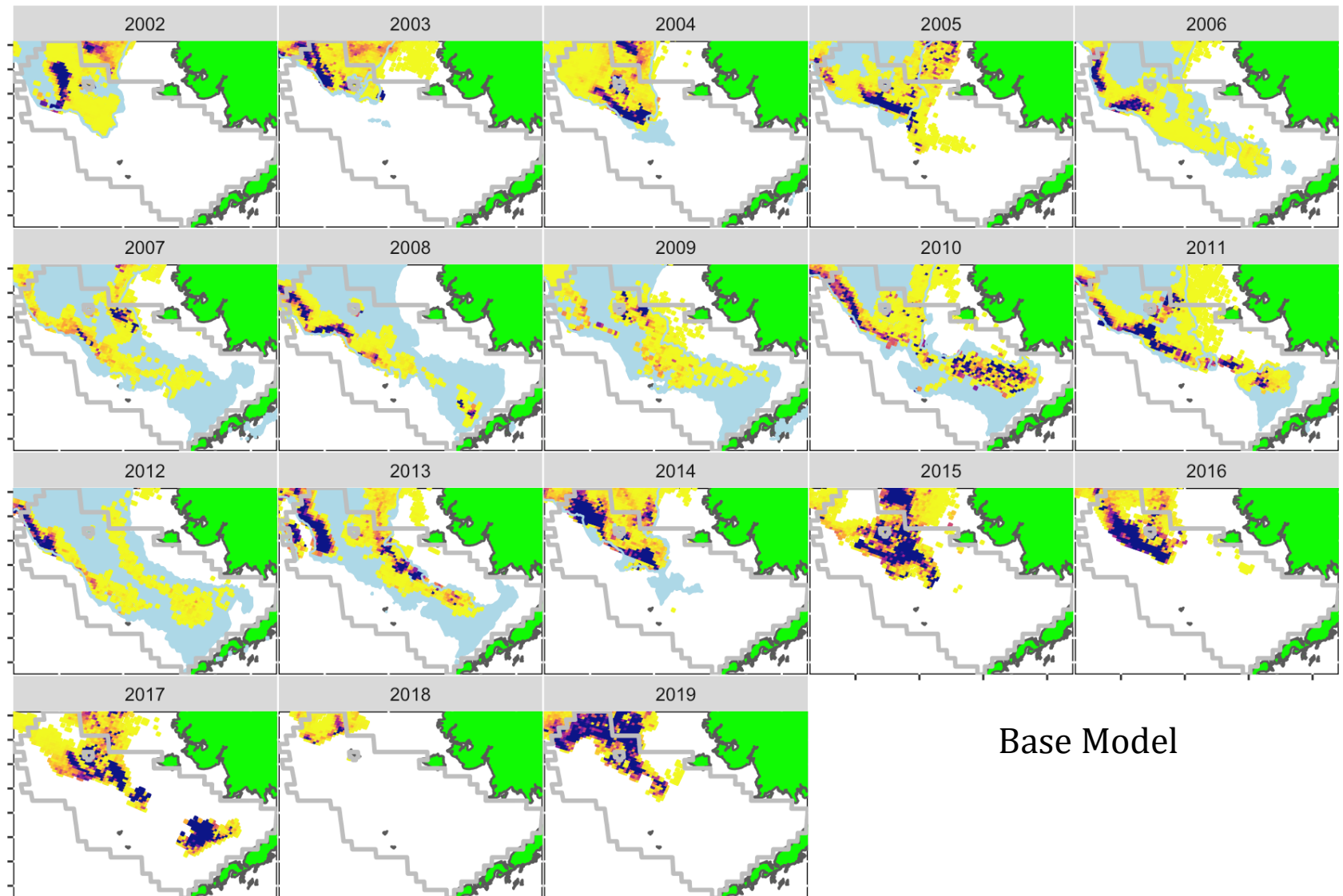
Base Model



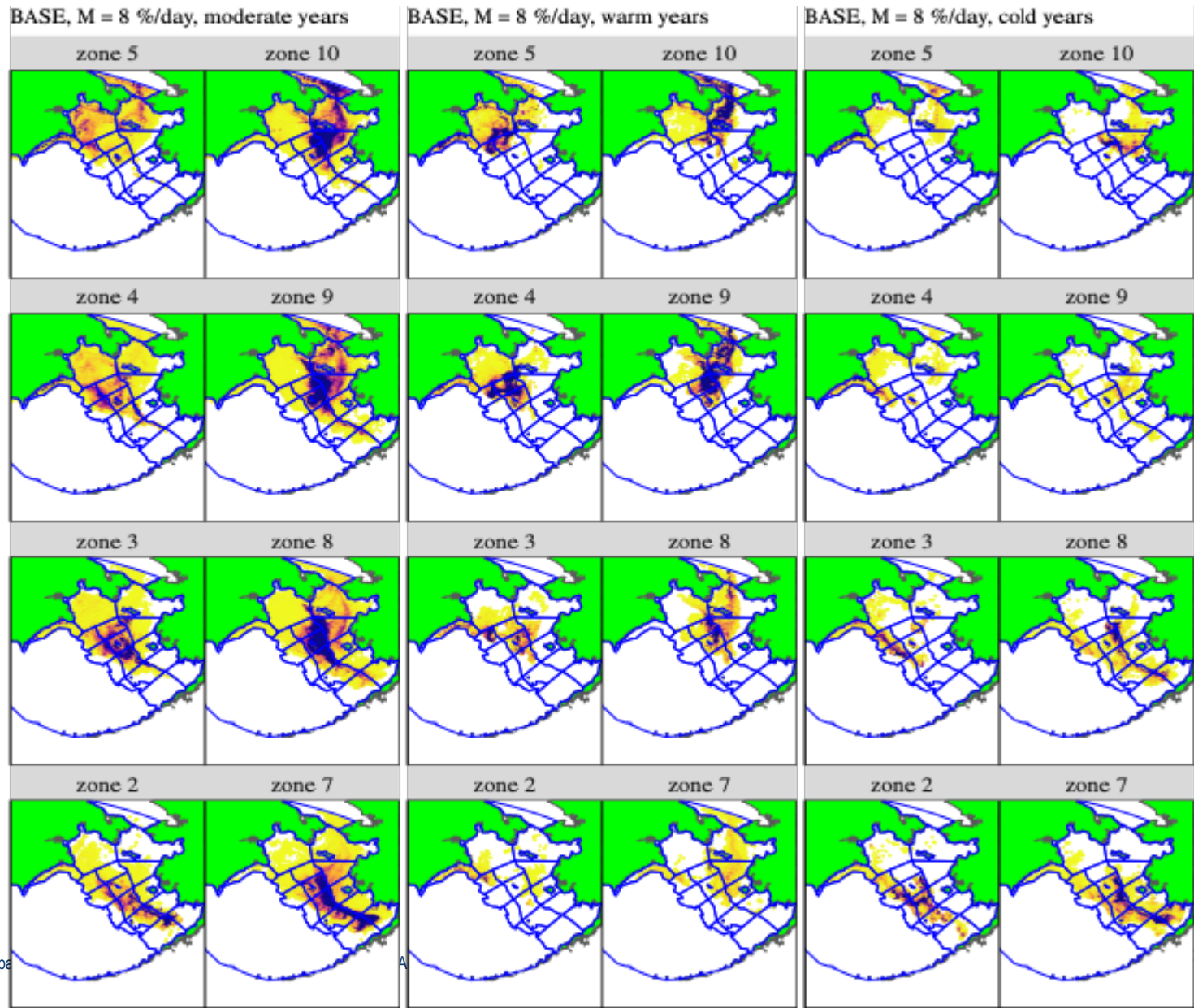
# Settlement Patterns: 2002-2019

assumed: 8%/day mortality

Base Model

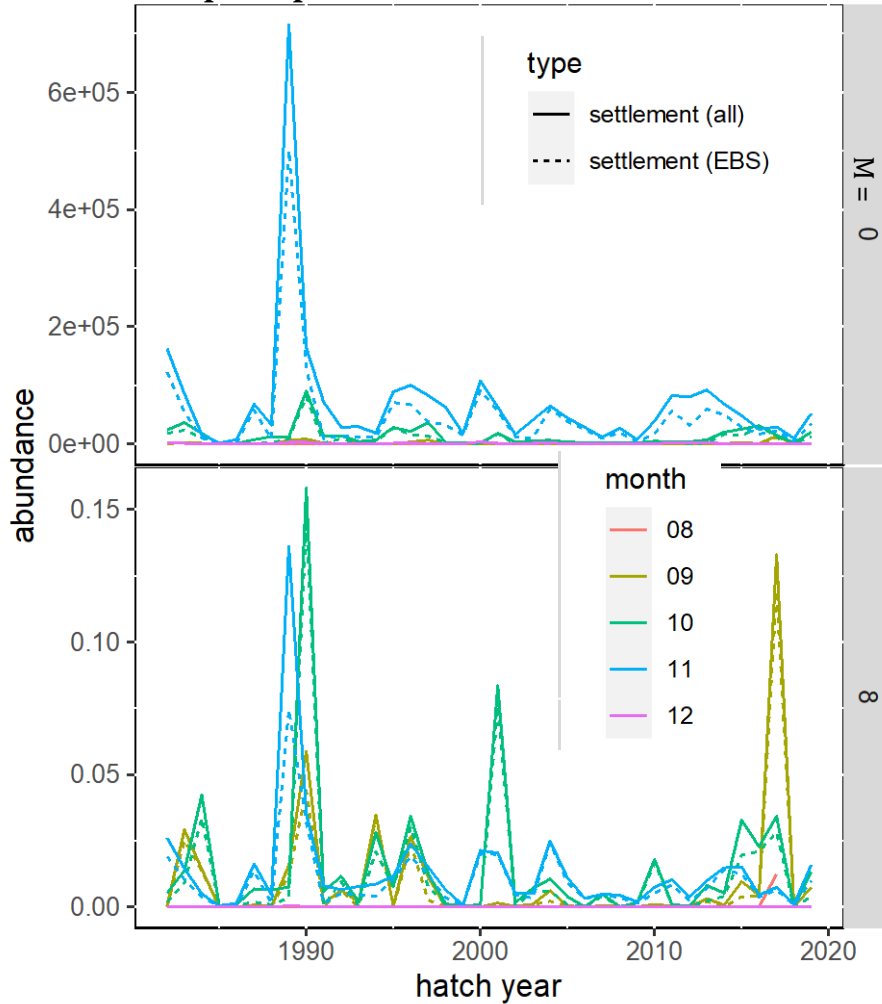


# Settlement results by temperature regime

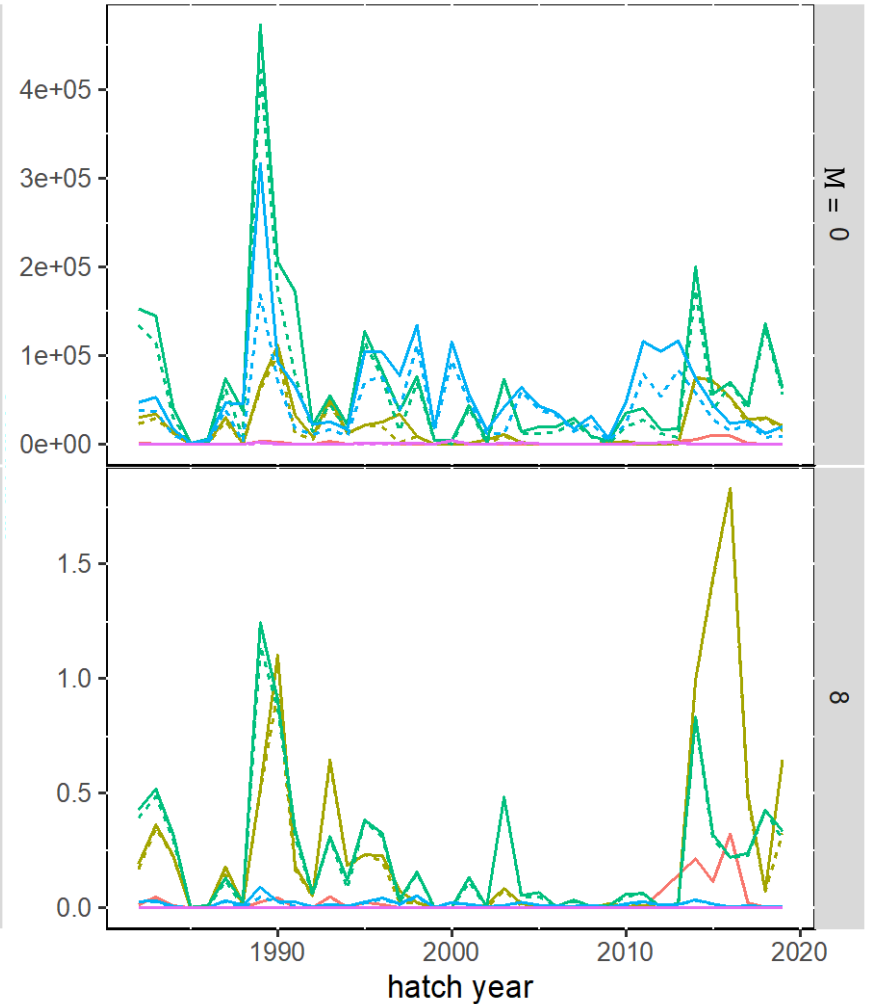


# Settlement Abundance Time Series

## temp-dependent molt duration



## fixed molt duration





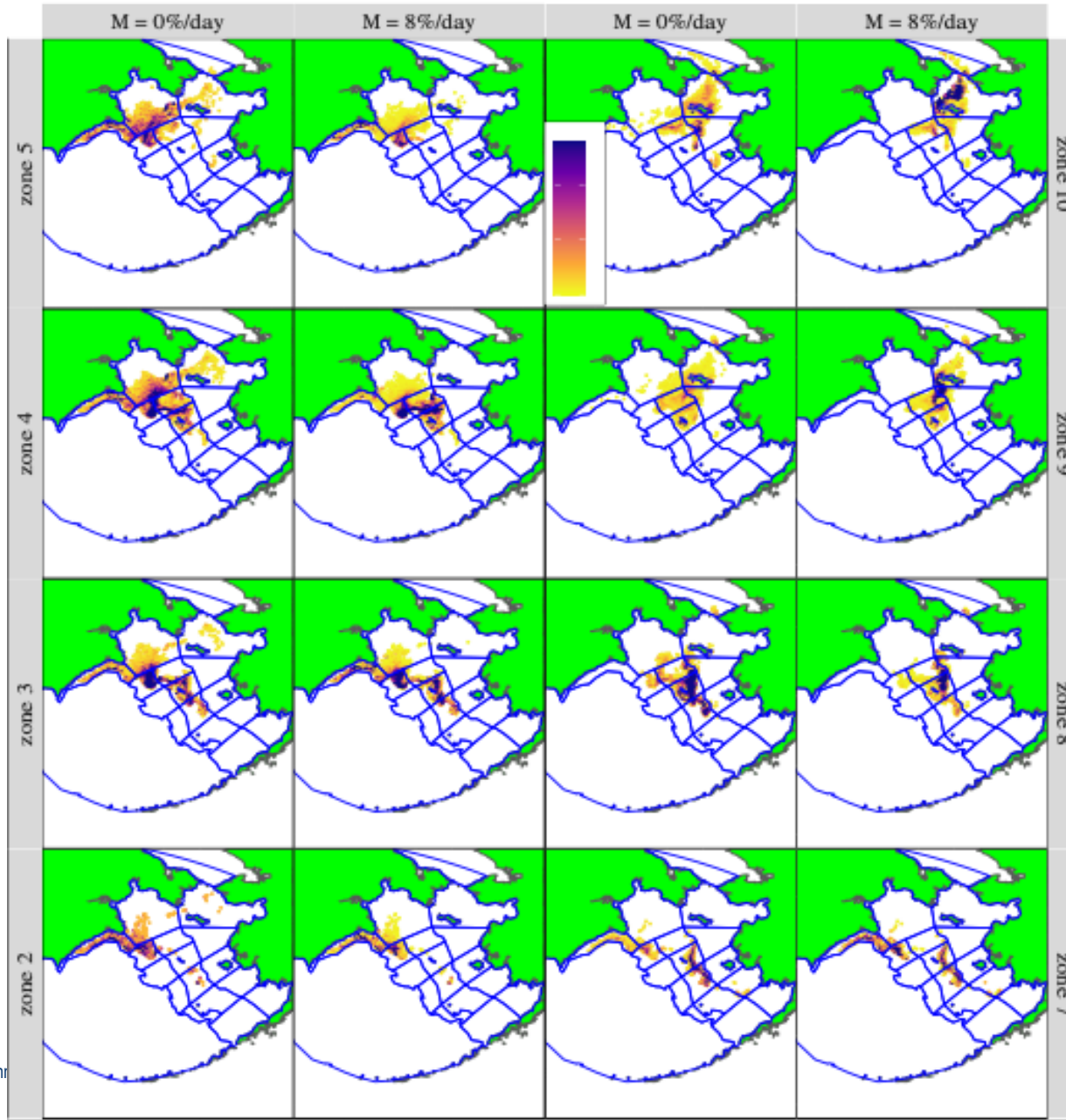
# Model uncertainty

- within-season hatch variation
- benthic nursery characteristics
  - temperature range
  - bottom type
  - other factors
- mortality rates
  - starvation
  - predation



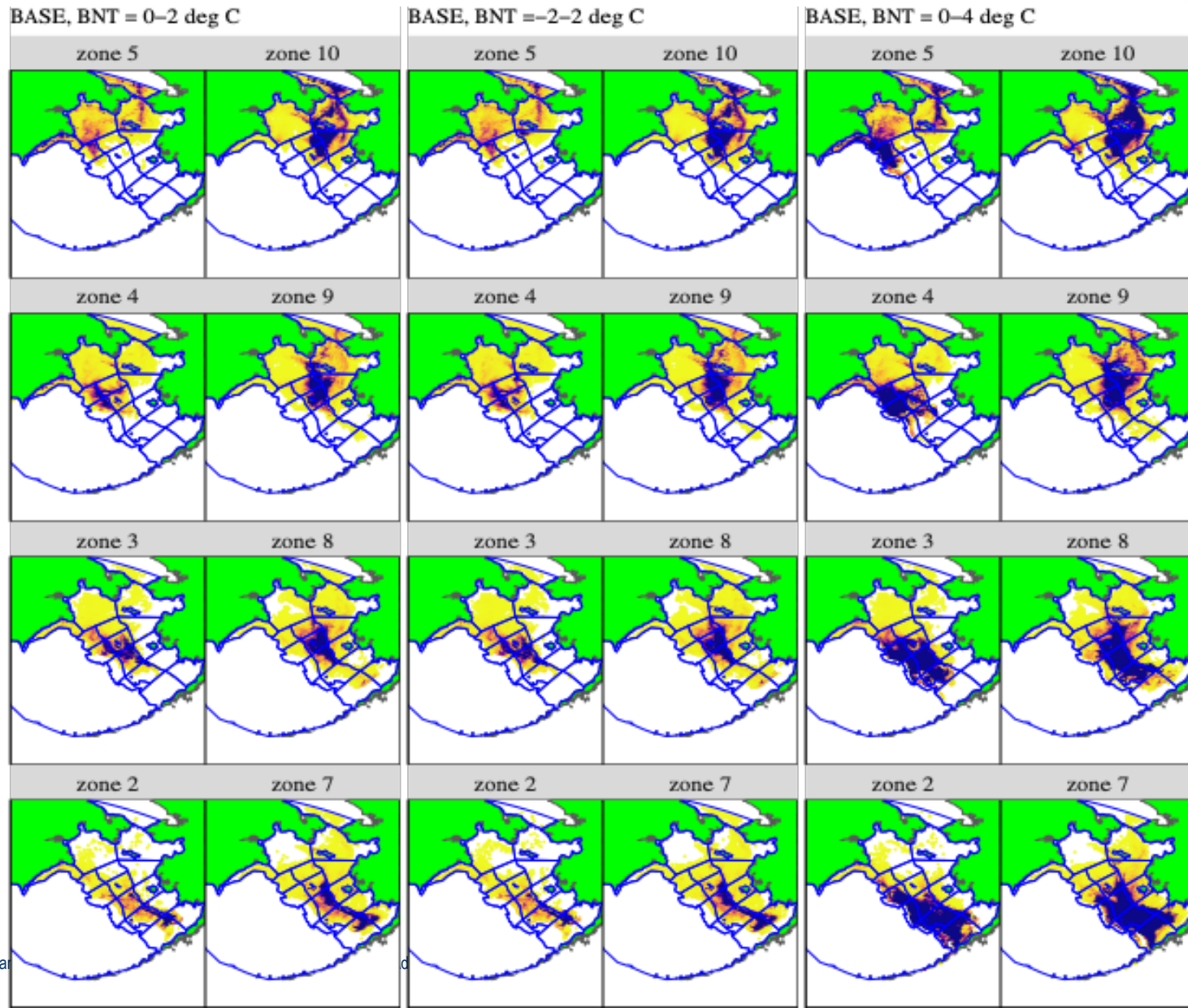
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# Model uncertainty: Mortality rates

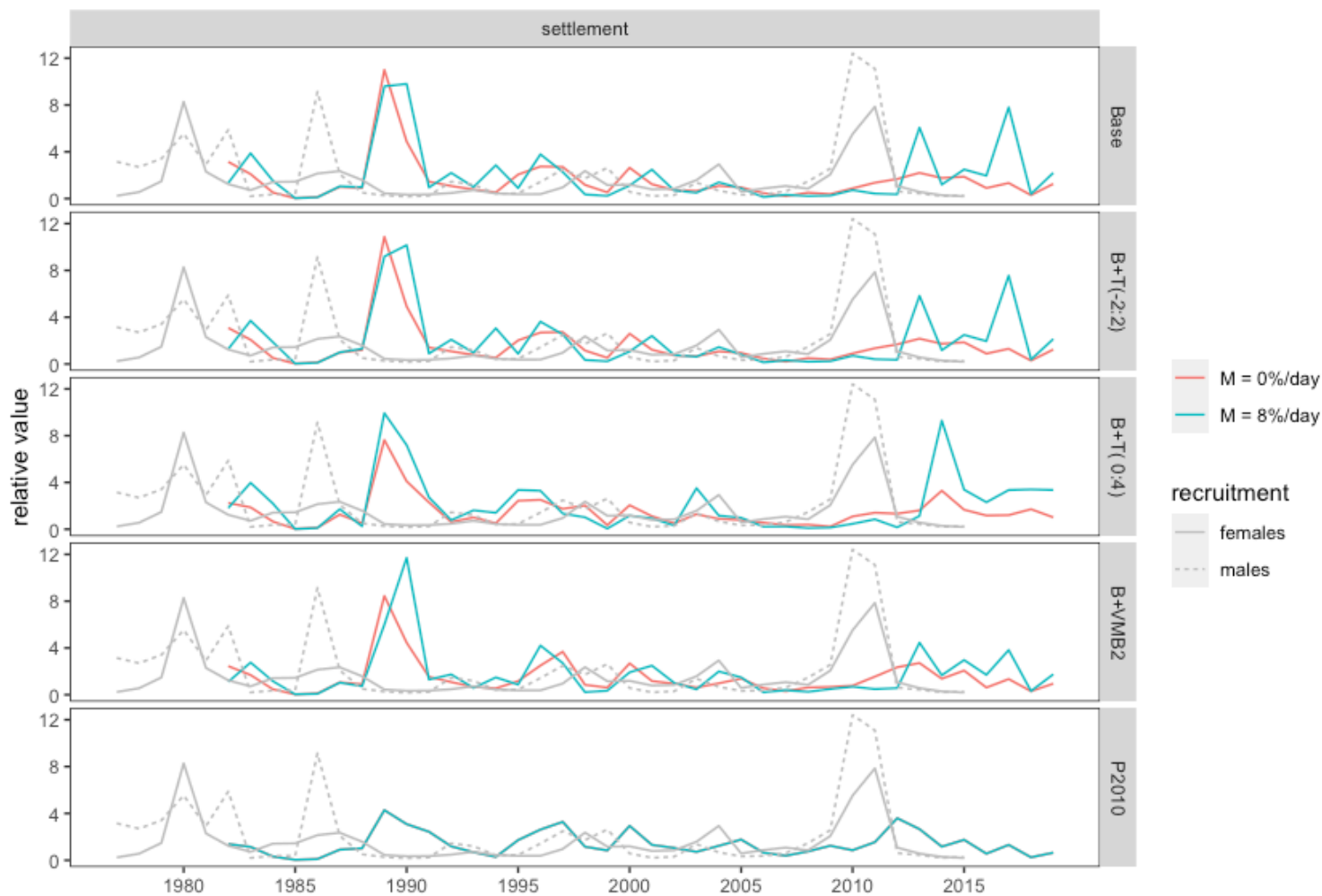


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# Model uncertainty: variability with BNT



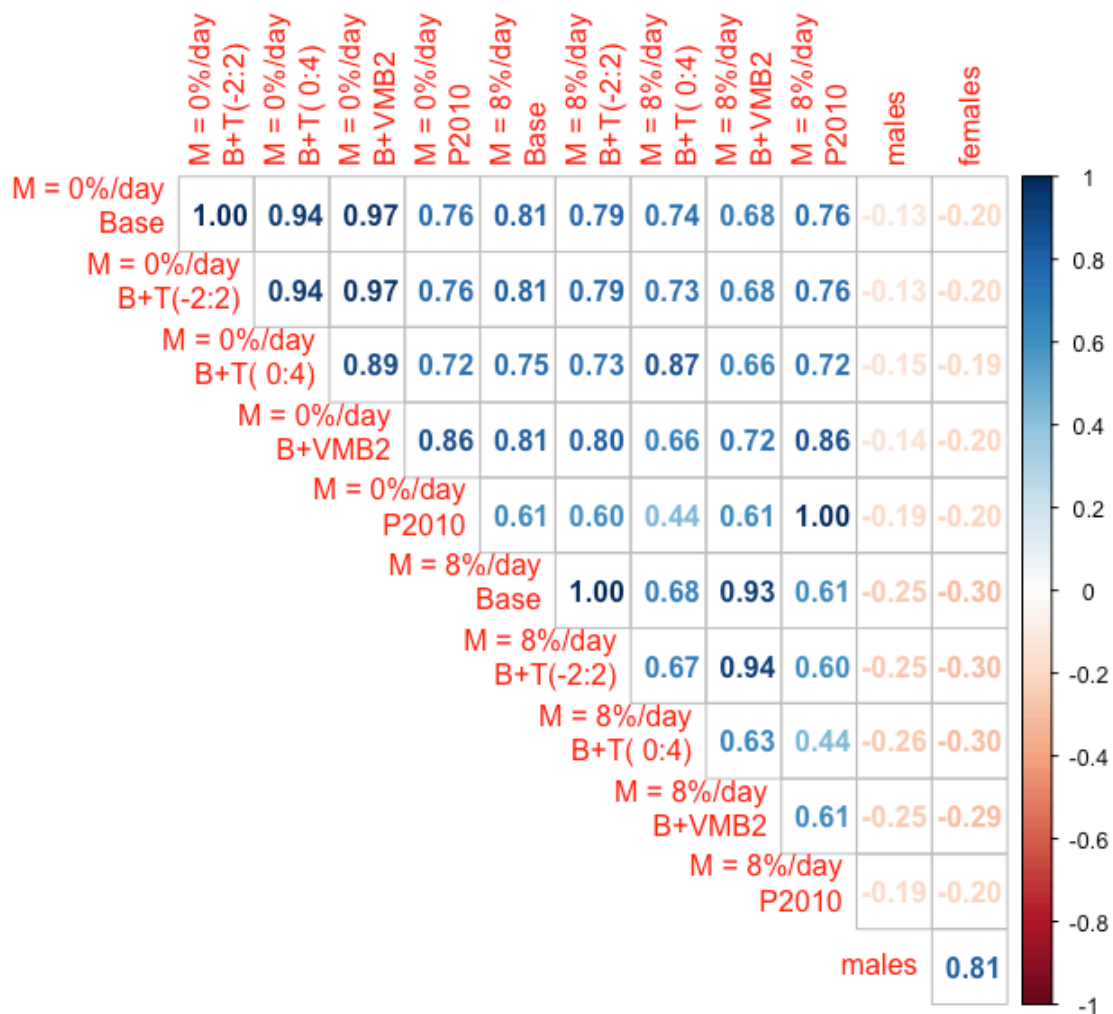
# Predicted Settlement And Recruitment



Recruitment lagged 5 years



# Correlations between settlement and recruitment



Recruitment lagged 5 years

# IBM Results: Key points

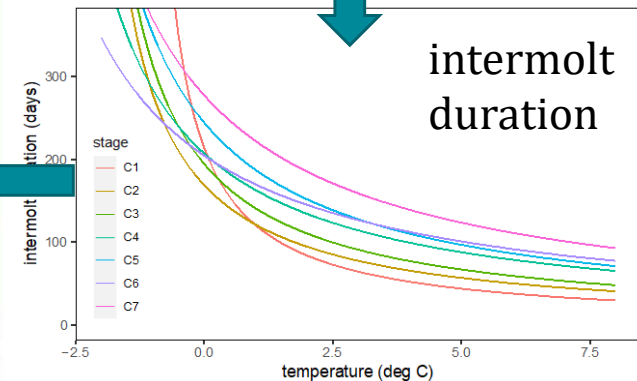
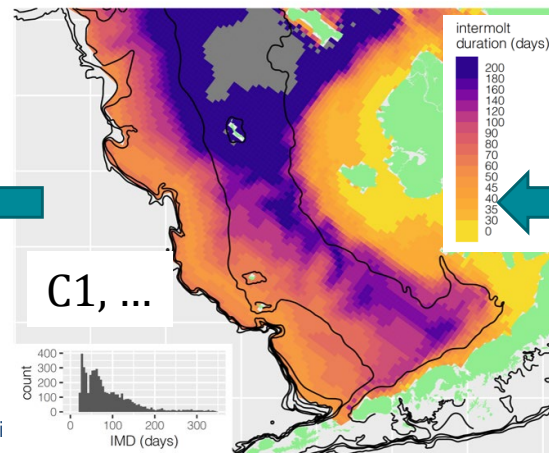
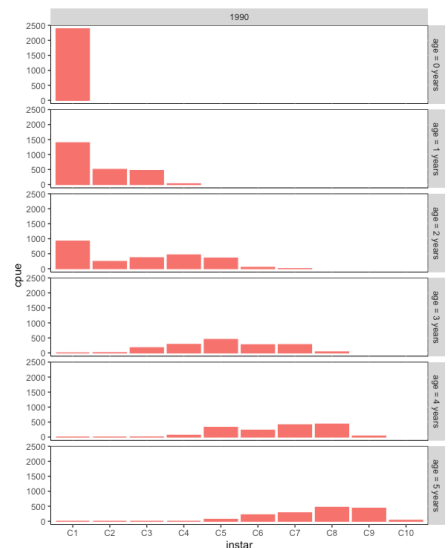
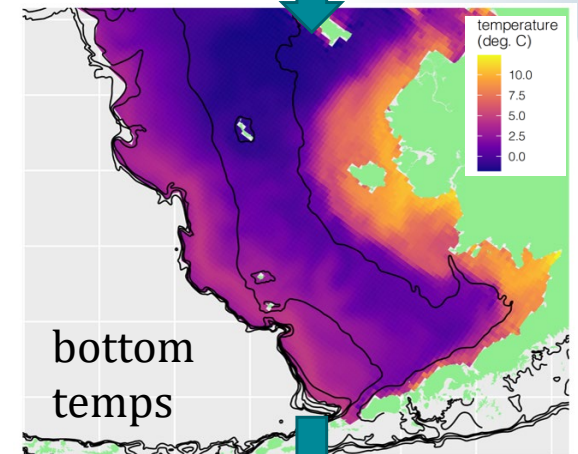
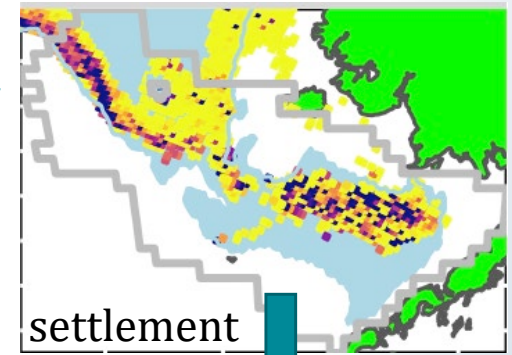
- integrates what we (think we) know about early life stages
  - lots we don't know
- EBS likely contributes to Chukchi stocks
- Some transport to southeast (counter to ratchet effect)
- moderate temperature years may be most productive
  - warmer temps -> faster development -> less pelagic mortality
  - colder temps -> more extensive settlement habitat
- settlement and recruitment are decoupled:
  - predicted settlement abundance has little relationship to estimated recruitment 5 years in the future
  - possibilities
    - IBM is missing a key factor?
      - starvation? spatiotemporal patterns in predation?
    - decoupling due to processes in benthic nurseries?



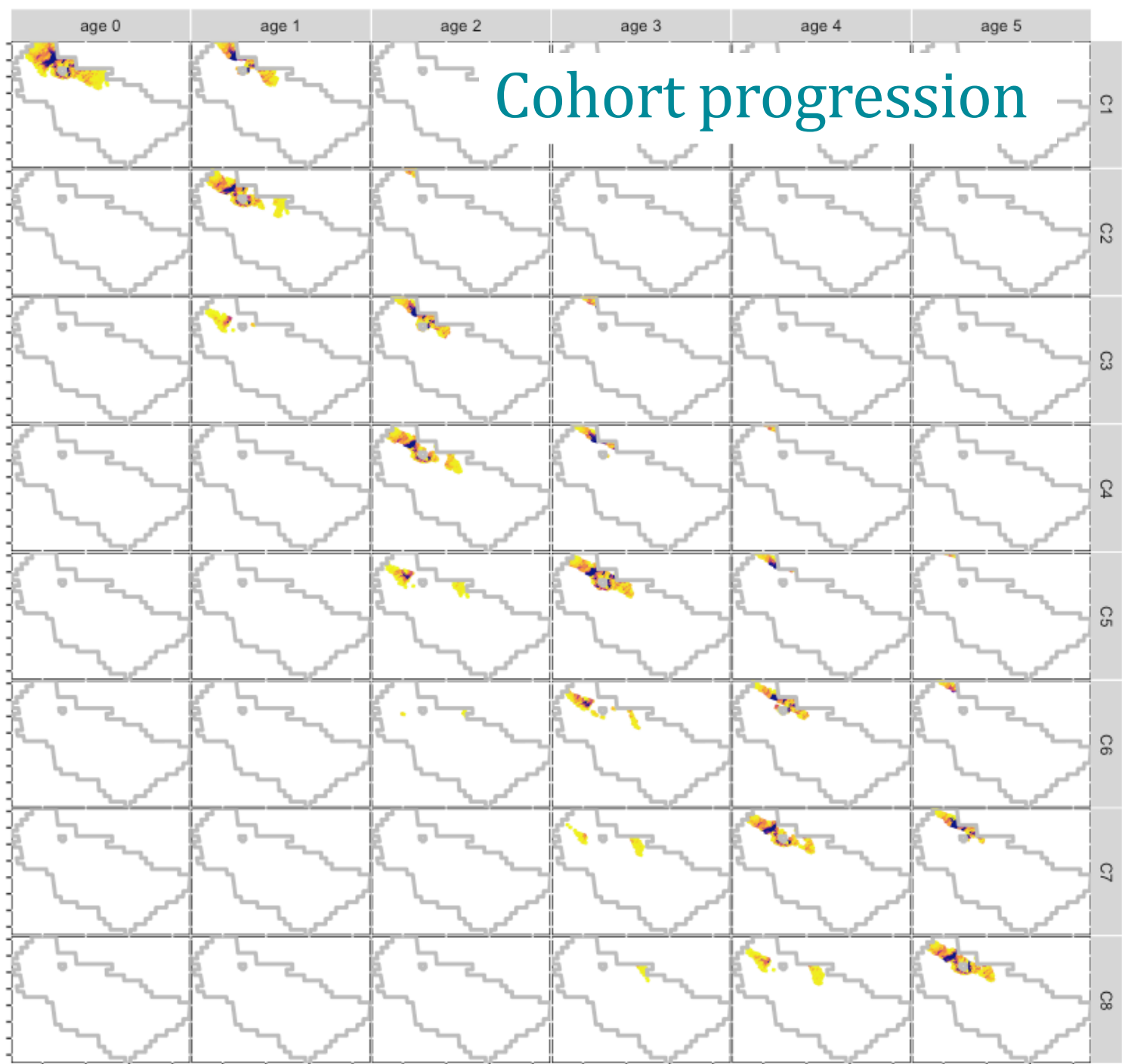
# Benthic Cohort Progression Model

- Initial pattern: IBM settlement pattern by ROMS grid cell
- Molt stage projected using ROMS bottom temps, lab-determined development rates\*
- Can include instar-specific, temperature-dependent mortality rates (“cod predation”)
- Daily integration for abundance, instar within each ROMS grid cell

\*Yamamoto et al (2015)



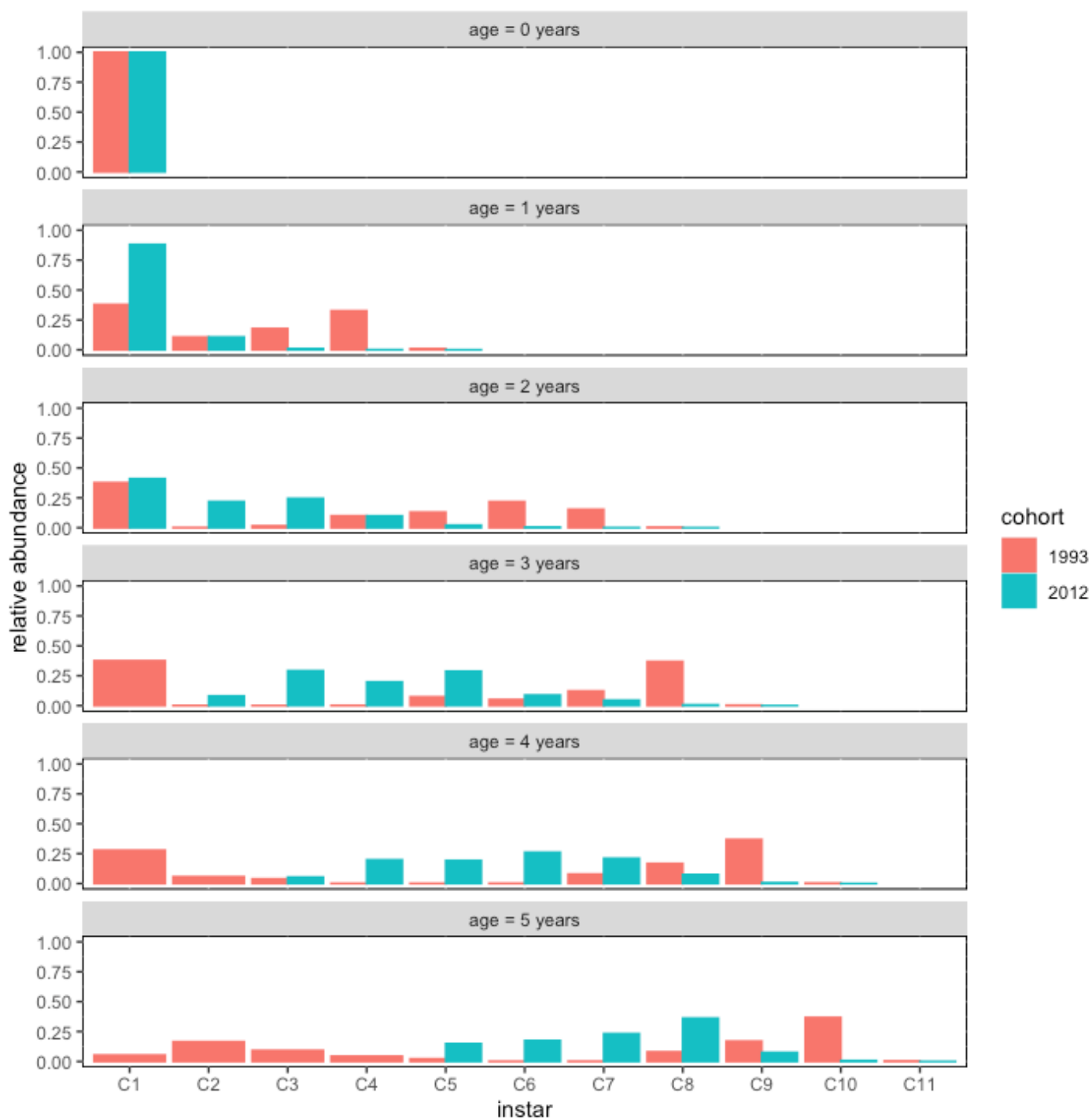
# Cohort progression





# Cohort Progression: Warm vs. Cold Years

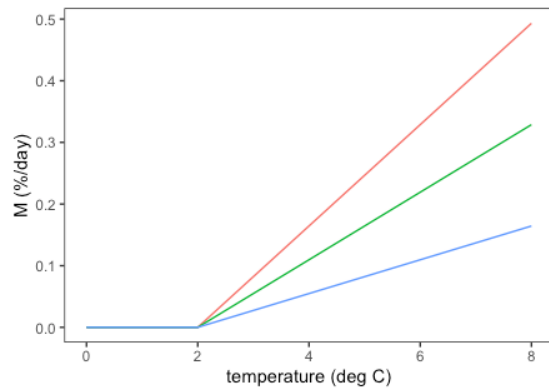
- no mortality



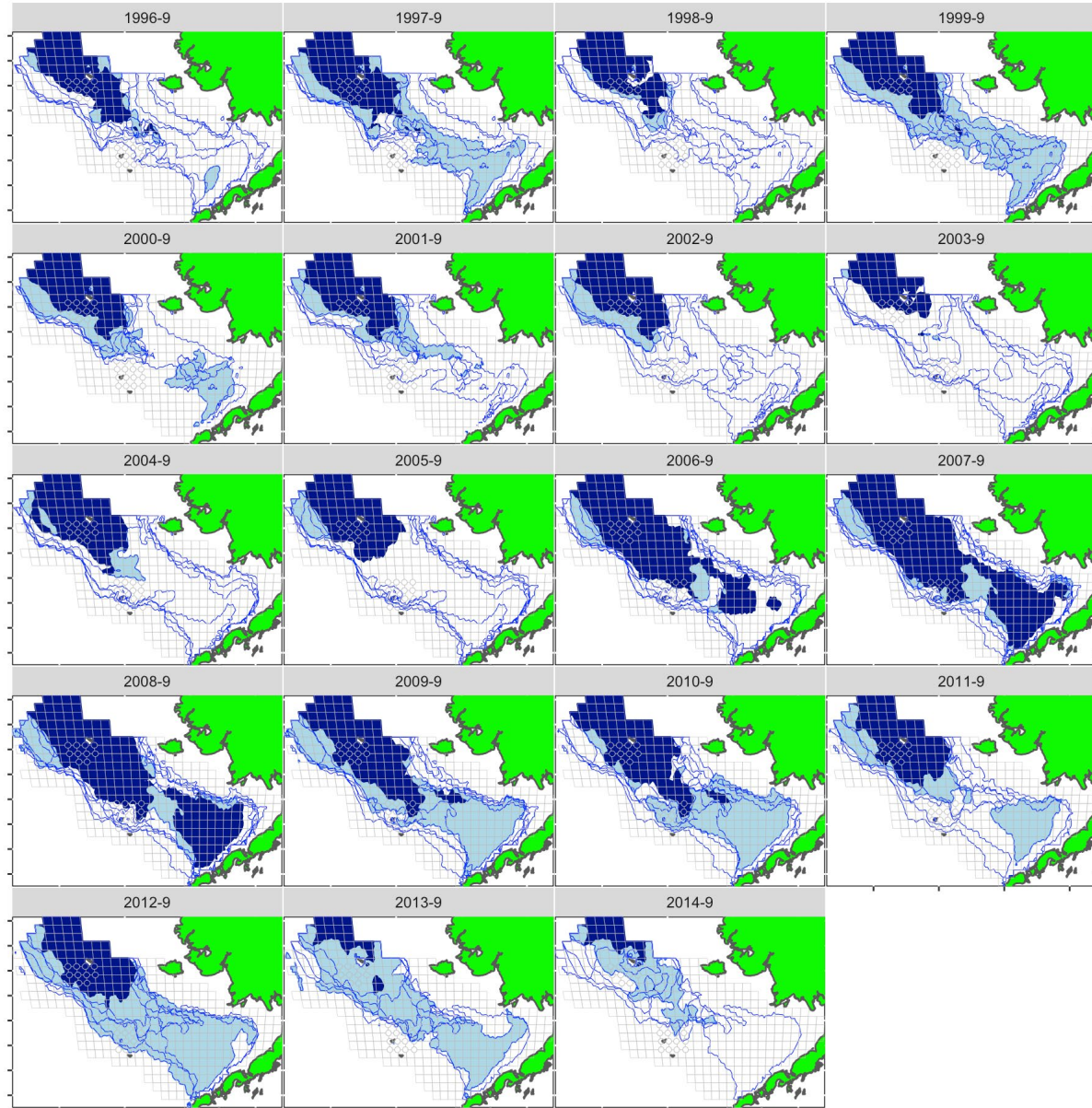
# Cold pool extents: settlement to recruitment

by settlement year

- predation by cod excluded from cold pools?



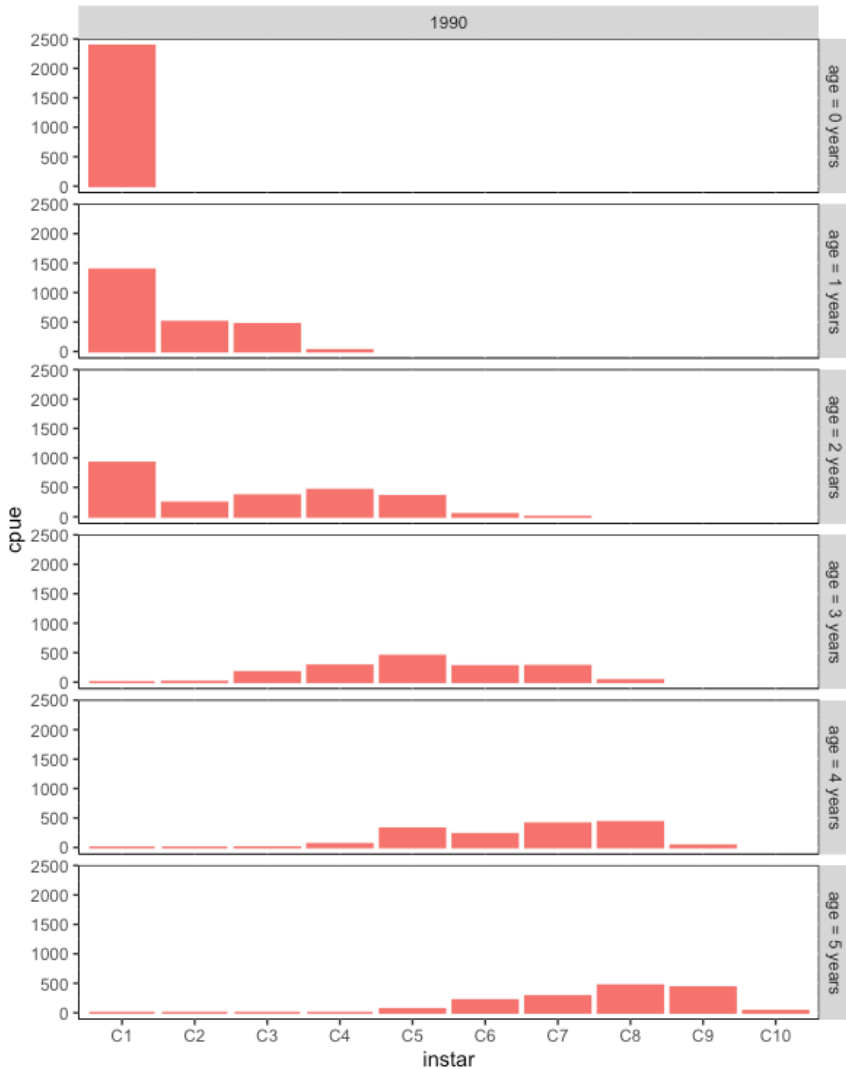
instar  
— C1  
— C2  
— C3



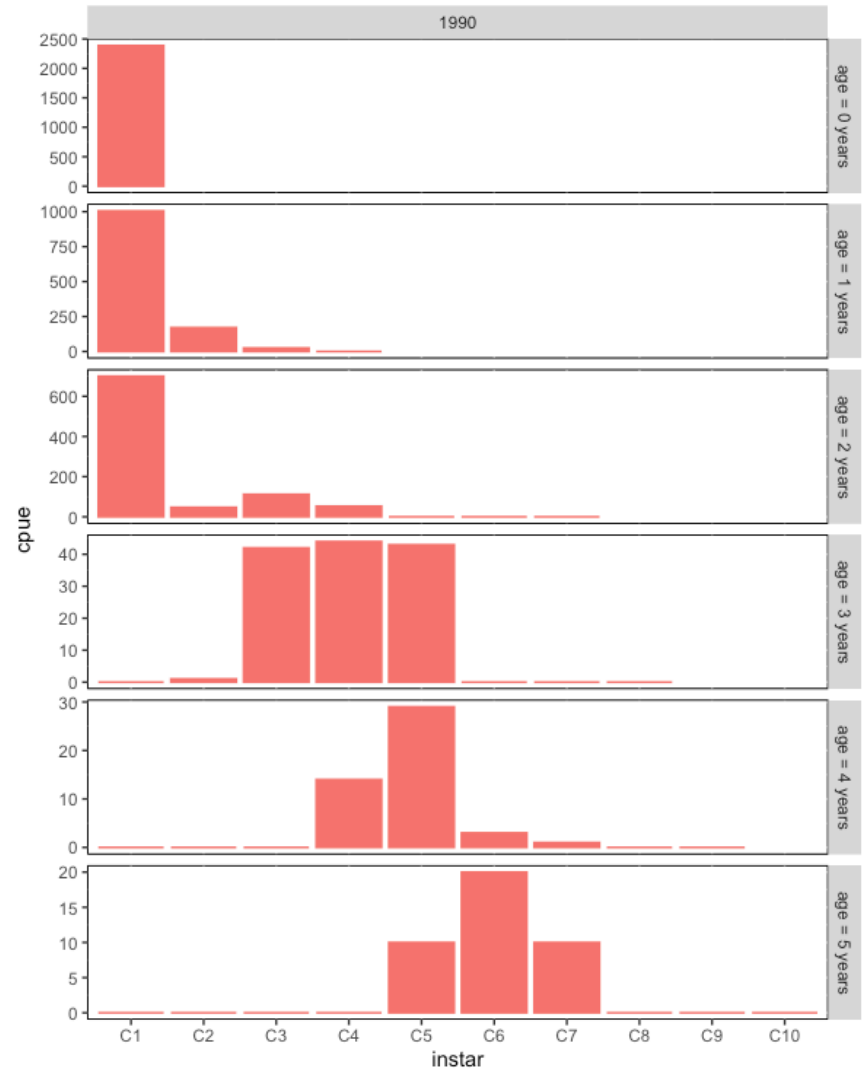
# Example Cohort Progression

- C5: 16 mm CW
- C6: 21 mm CW
- C7: 30 mm CW

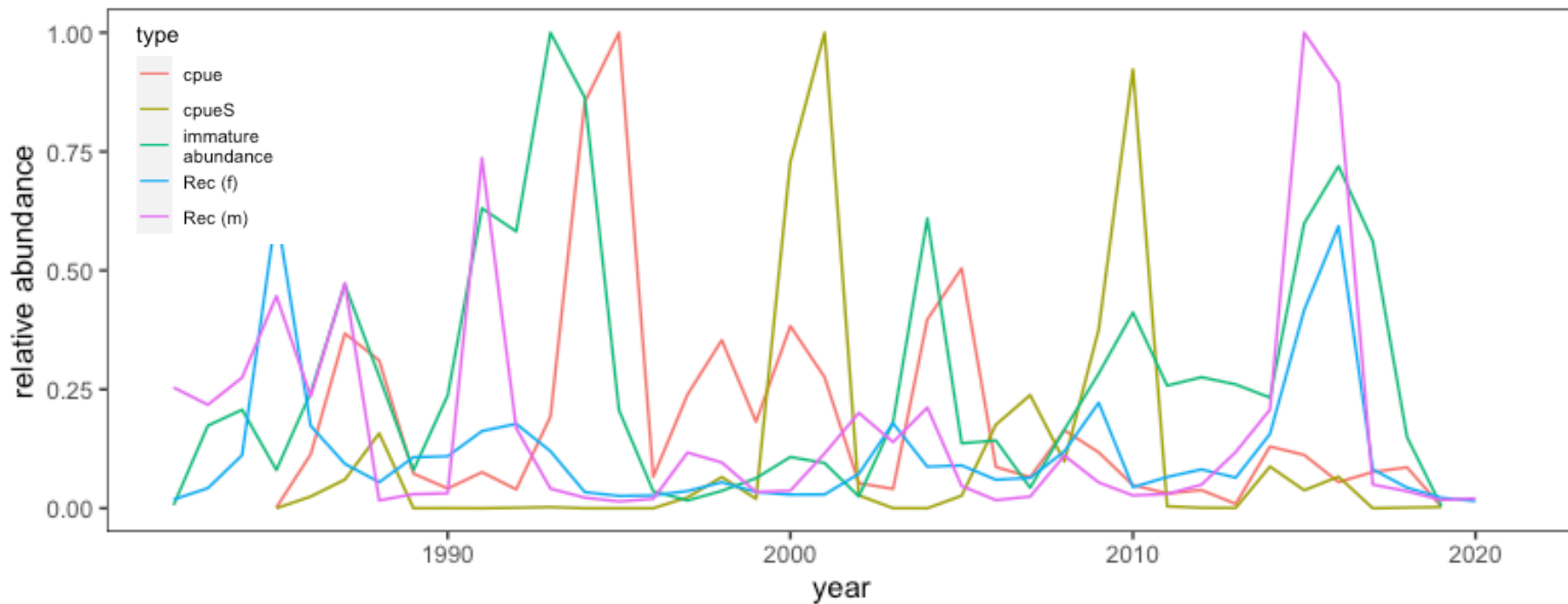
no "cod predation"



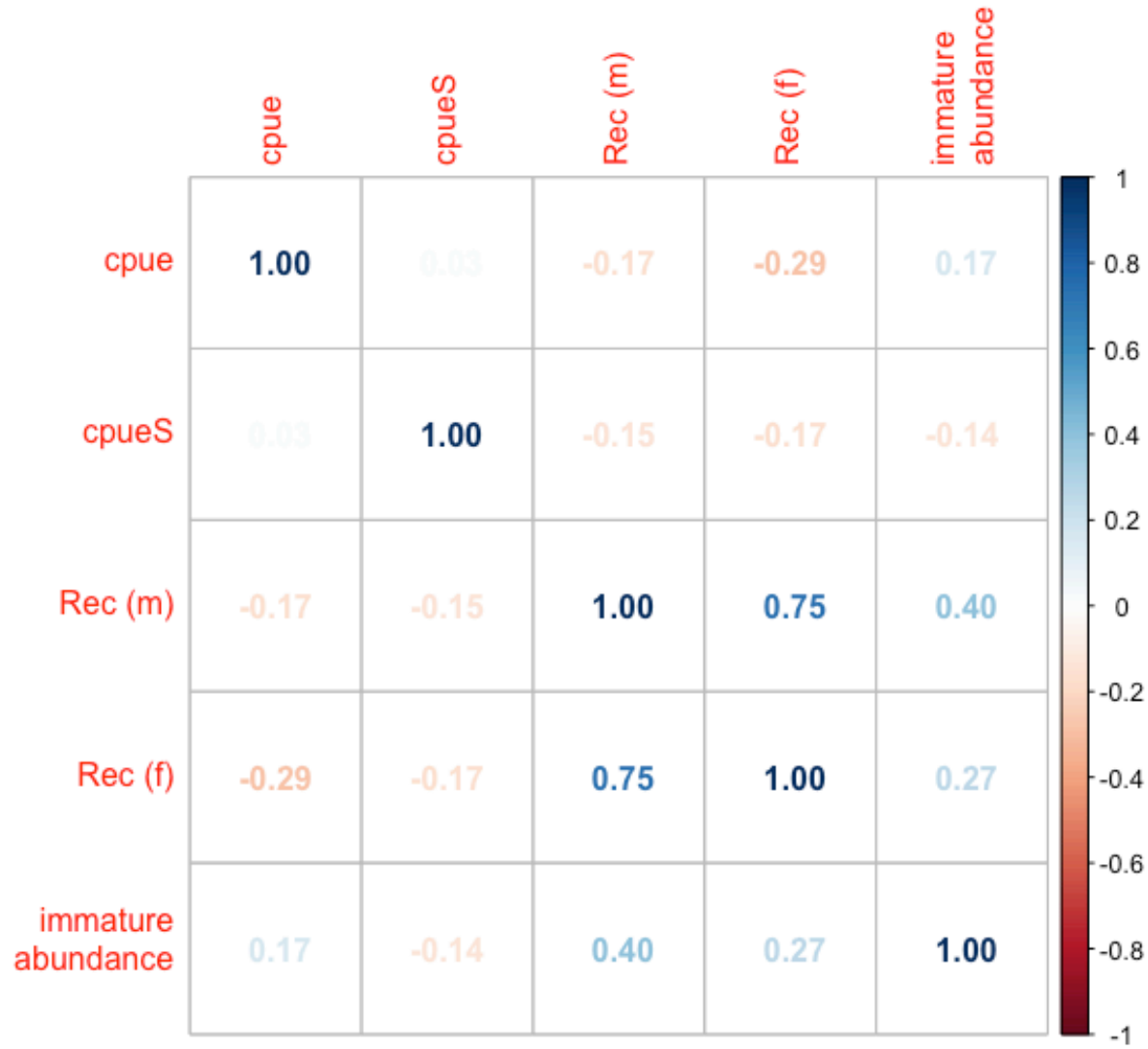
with "cod predation"



# Indices



# Indices



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# Summary

- Bit disappointing in terms of correlation with recruitment
- What's missing?
  - better description of P. cod predation?
  - density dependence?
  - other mortality factors?
- Next steps?



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