

Climate forcing of *C. finmarchicus* populations of the North Atlantic

WHOI

McGillicuddy

Smith

Wiebe

Copley

UConn

Bucklin

Unal

Rutgers

Haidvogel

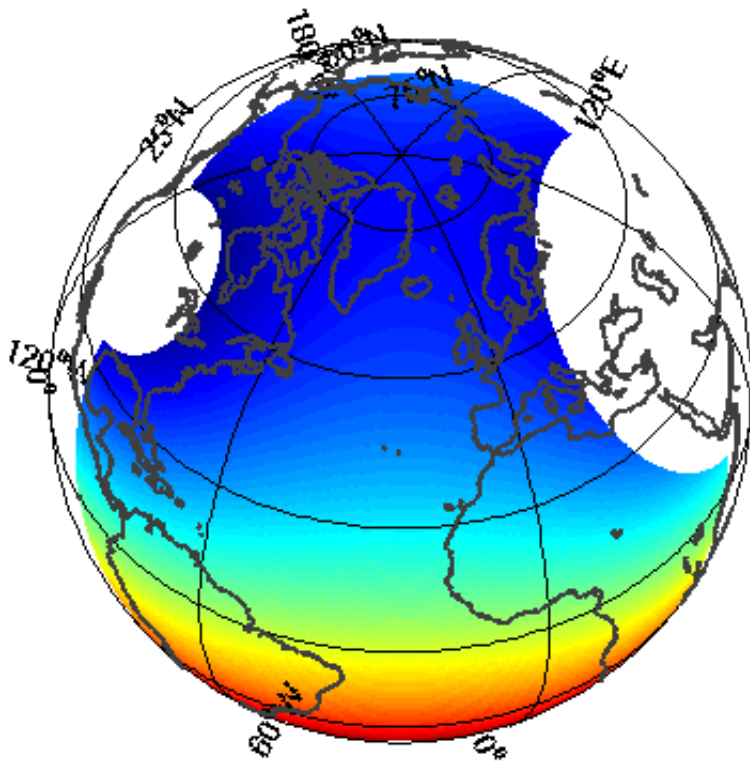
Levin

Overall Objectives

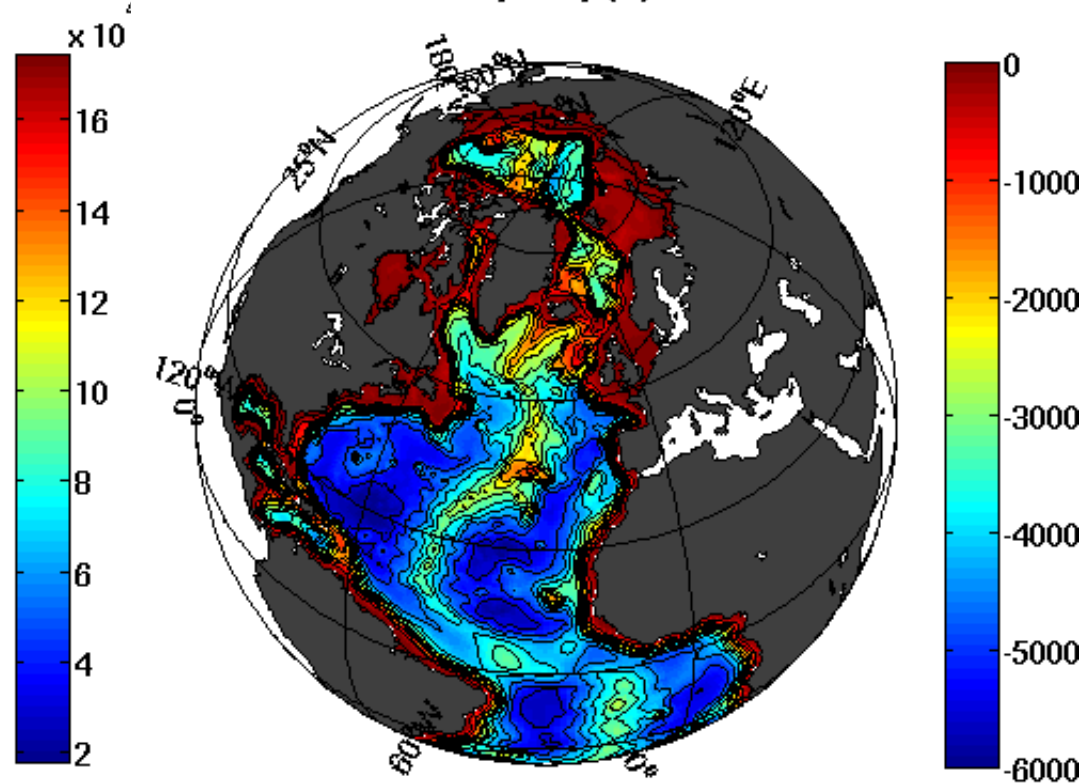
- Inversions of the “mean state”
 - Climatological mean seasonal forcing
 - Diapause entry hypotheses: food, photoperiod
 - Diapause exit hypotheses: development, photoperiod
 - Climatological CPR data
 - Control parameter: mortality (spatially variable, stage dependent)
 - Skill assessment: cross-validation
- Use the genetic data to estimate the rate of population exchange between gyres, and compare with model predictions of same
- Investigate interannual to decadal variability
 - High-NAO state vs. low-NAO state
 - Hindcast 1950s-present

Climatological low-resolution North Atlantic ROMS (NATL-LR) simulation

Grid Spacing (m)



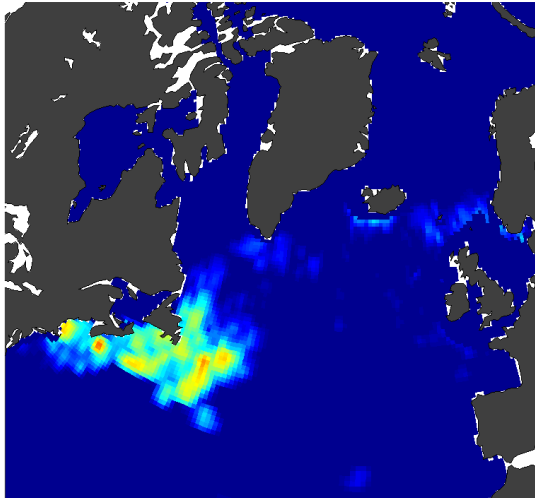
Bathymetry (m)



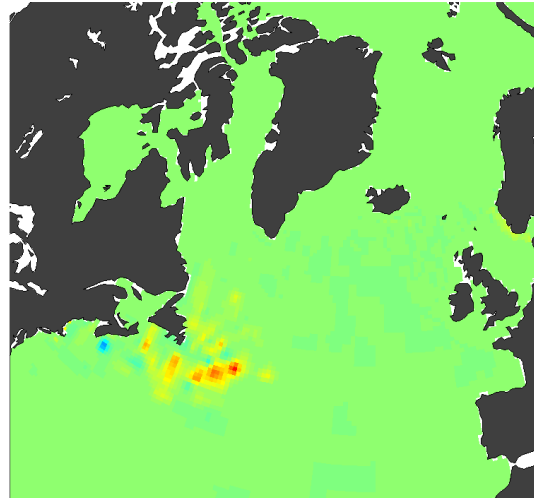
The grid is a modification the grid designed by Paul Budgell. The grid has 20 to 170 km resolution, region of interest has resolution within 40 km, 42 vertical layers.

Inverse solution for Jan-Feb (surface)

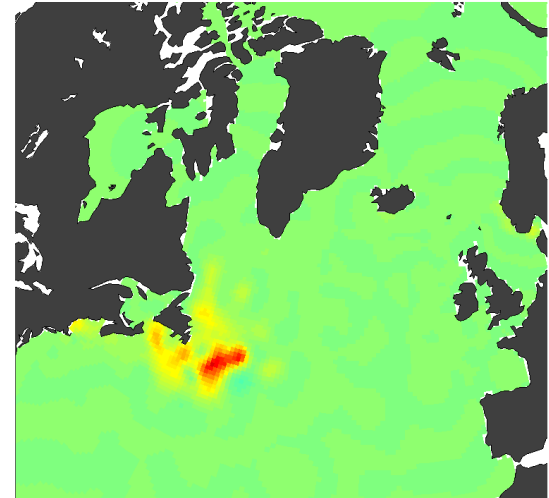
Initial condition



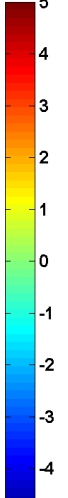
Poor man's source



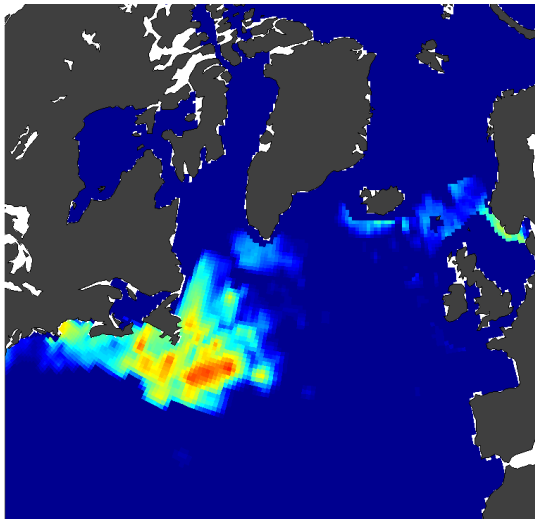
Inverse source



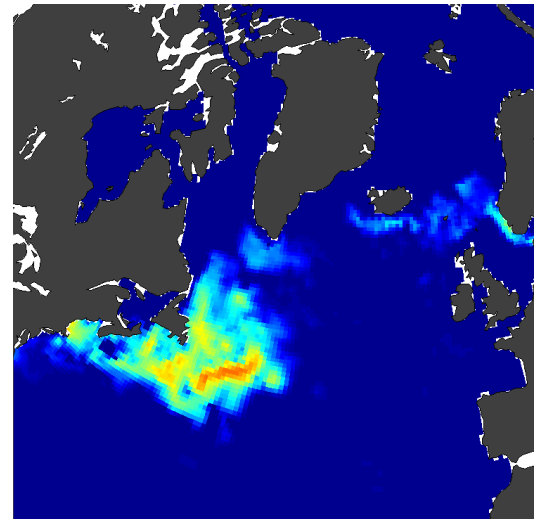
R: # m⁻³ d⁻¹



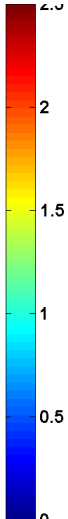
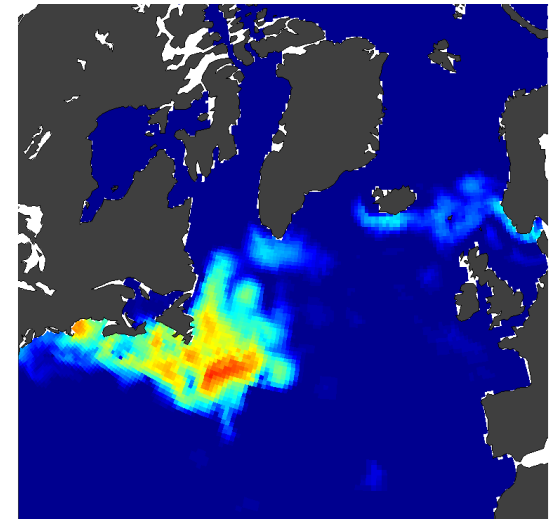
Data next month



with poor man's source



with inverse source



log₁₀(C+1)