

Linkages and connectivity of cold-water coral communities along the slope, canyons, and seamounts in the North Atlantic Ocean (an overview)

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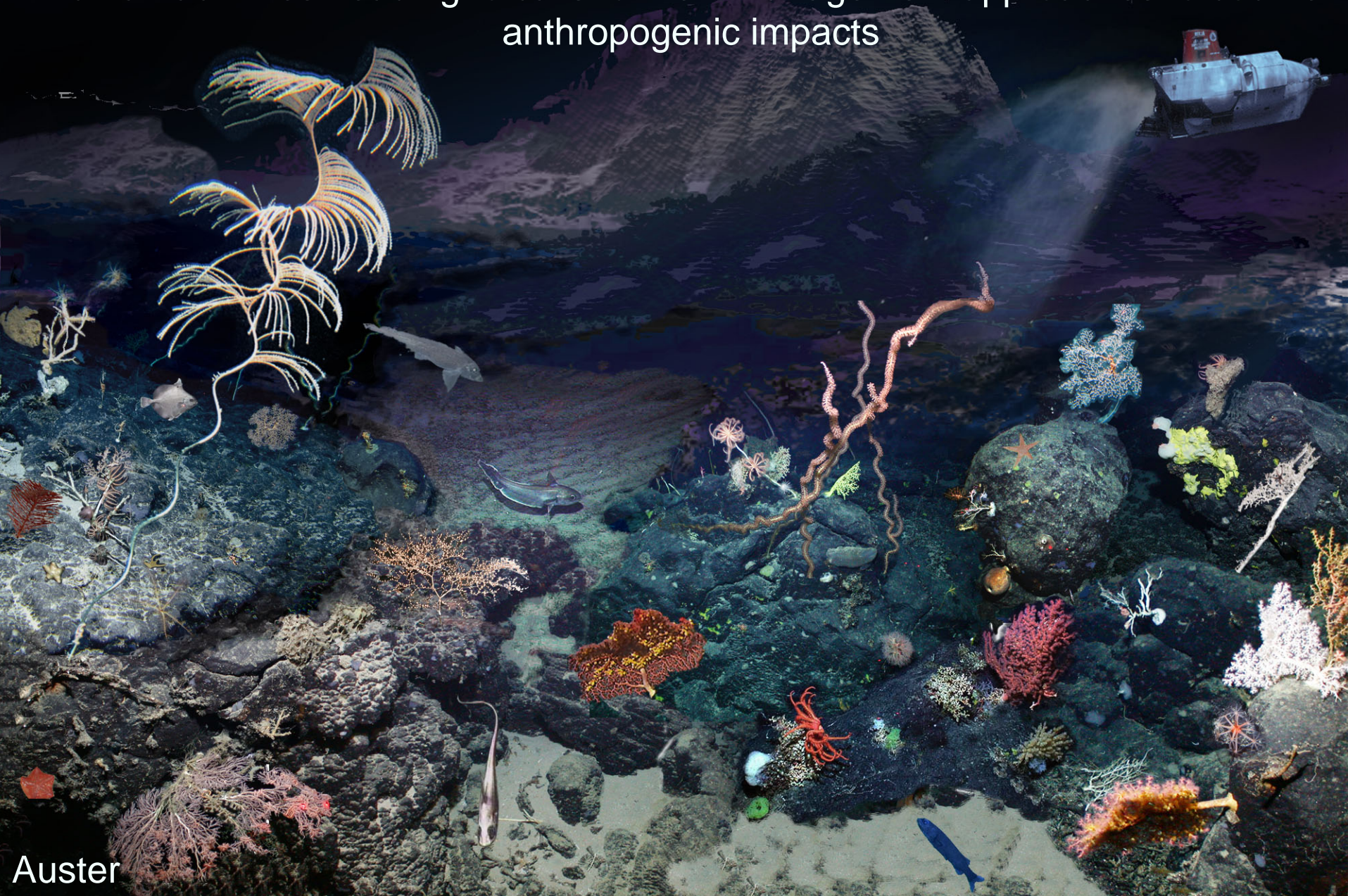
EXPLORE



TRACES: *Trans-Atlantic Coral Ecosystem Study*



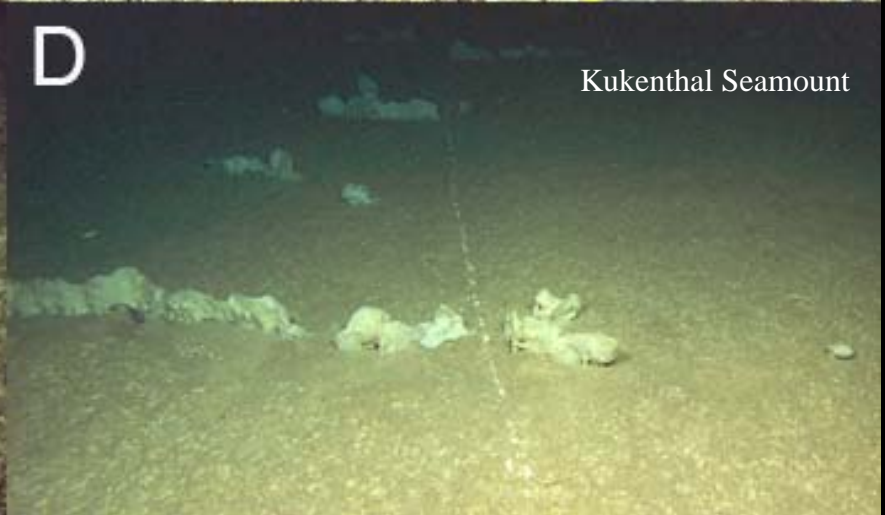
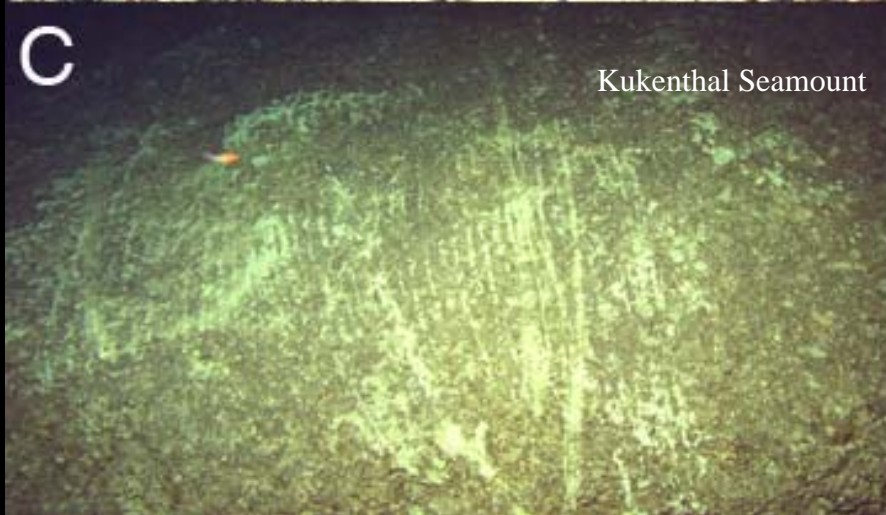
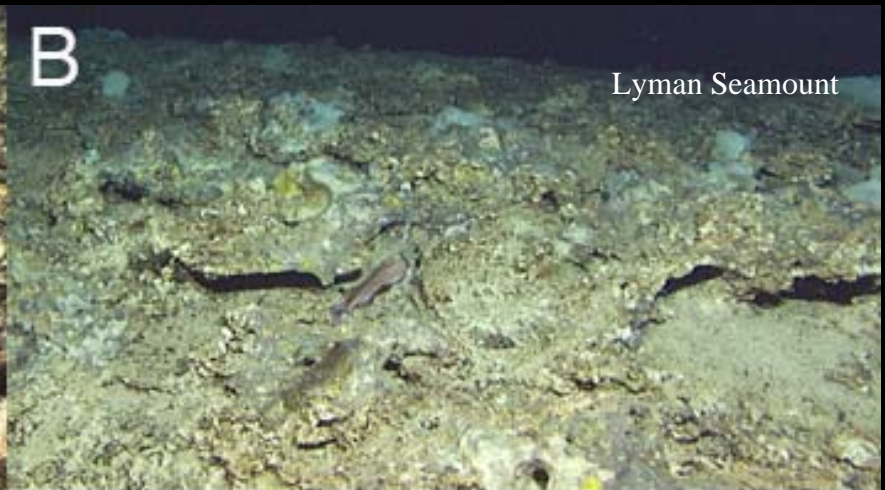
Coral ecosystems are under increasing and unprecedented stress from human activities- leading to calls for new management approaches to counter anthropogenic impacts



Coral Communities and Habitat= resources and consequences

- Low productivity and and slow recovery rates make coral ecosystems vulnerable to disturbance (e.g., fisheries activities and mining)





“Fisheries-related habitat damage” on the Corner Rise Seamounts (North Atlantic)

Exploitation (commercial recovery of minerals and fisheries):

- loss of habitat, prohibiting connectivity and reducing genetic diversity
- local, regional, or global extinction of endemic or rare taxa
- decreased diversity (all levels: genetic, species, phylogenetic, habitat)

Seamounts

DEPTH

- < 2000m
- > 2000m

Northwest Atlantic



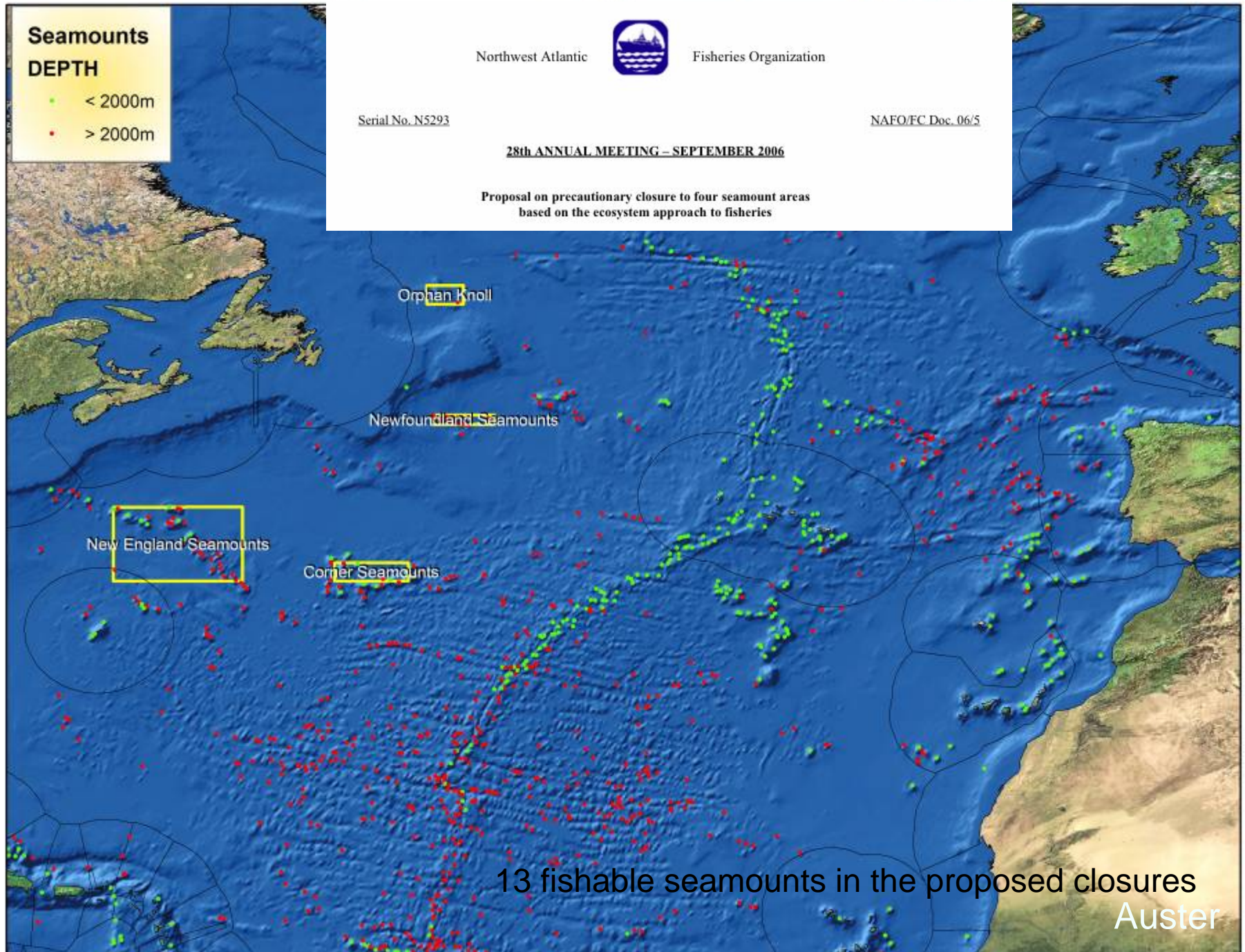
Fisheries Organization

Serial No. N5293

NAFO/FC Doc. 06/5

28th ANNUAL MEETING – SEPTEMBER 2006

**Proposal on precautionary closure to four seamount areas
based on the ecosystem approach to fisheries**



Policy Relevance and Drivers for Connectivity

“Connectivity plays the fundamental role in local and metapopulation dynamics, community dynamics and structure, genetic diversity and evolution, and resiliency of populations to human exploitation.

-Hastings and Harrison, 1994

“Understanding the ‘connectivity’ of marine populations is vital for conservation and fisheries management...the problem to be solved is the movement (and the controlling processes) of individuals and their contribution to the gene pool.

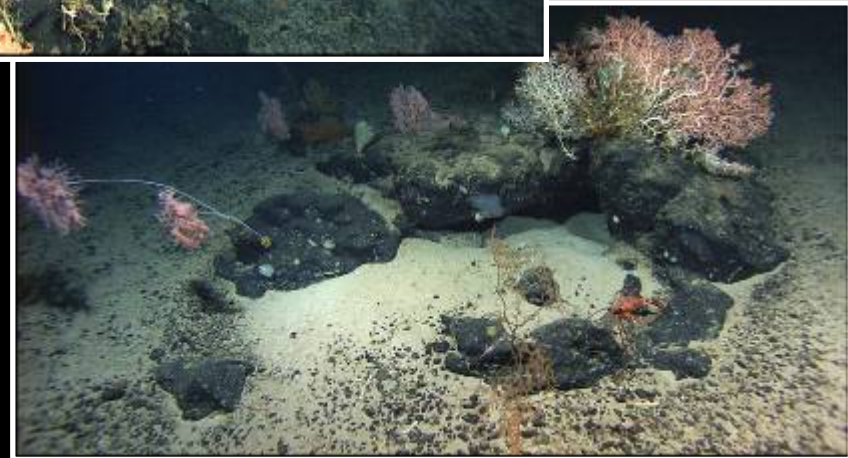
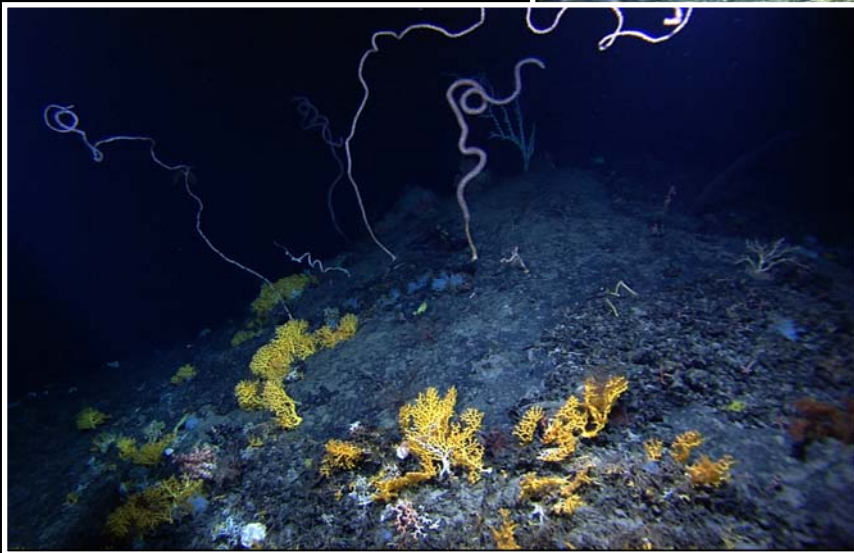
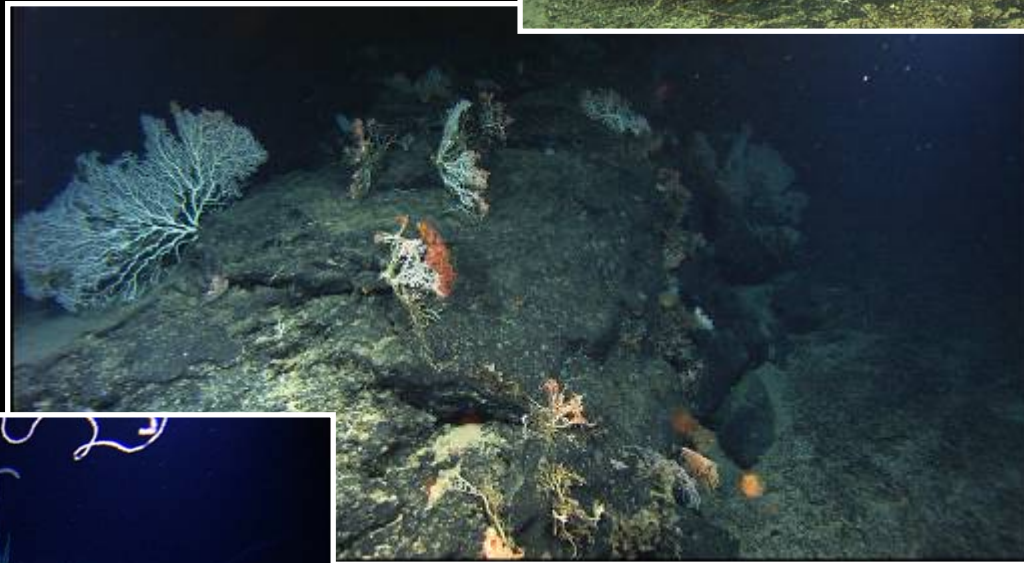
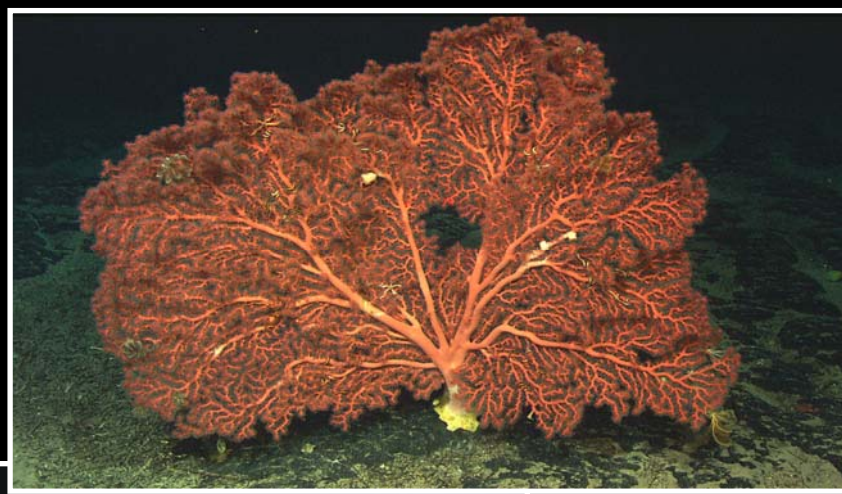
- Cowen et al., 2007

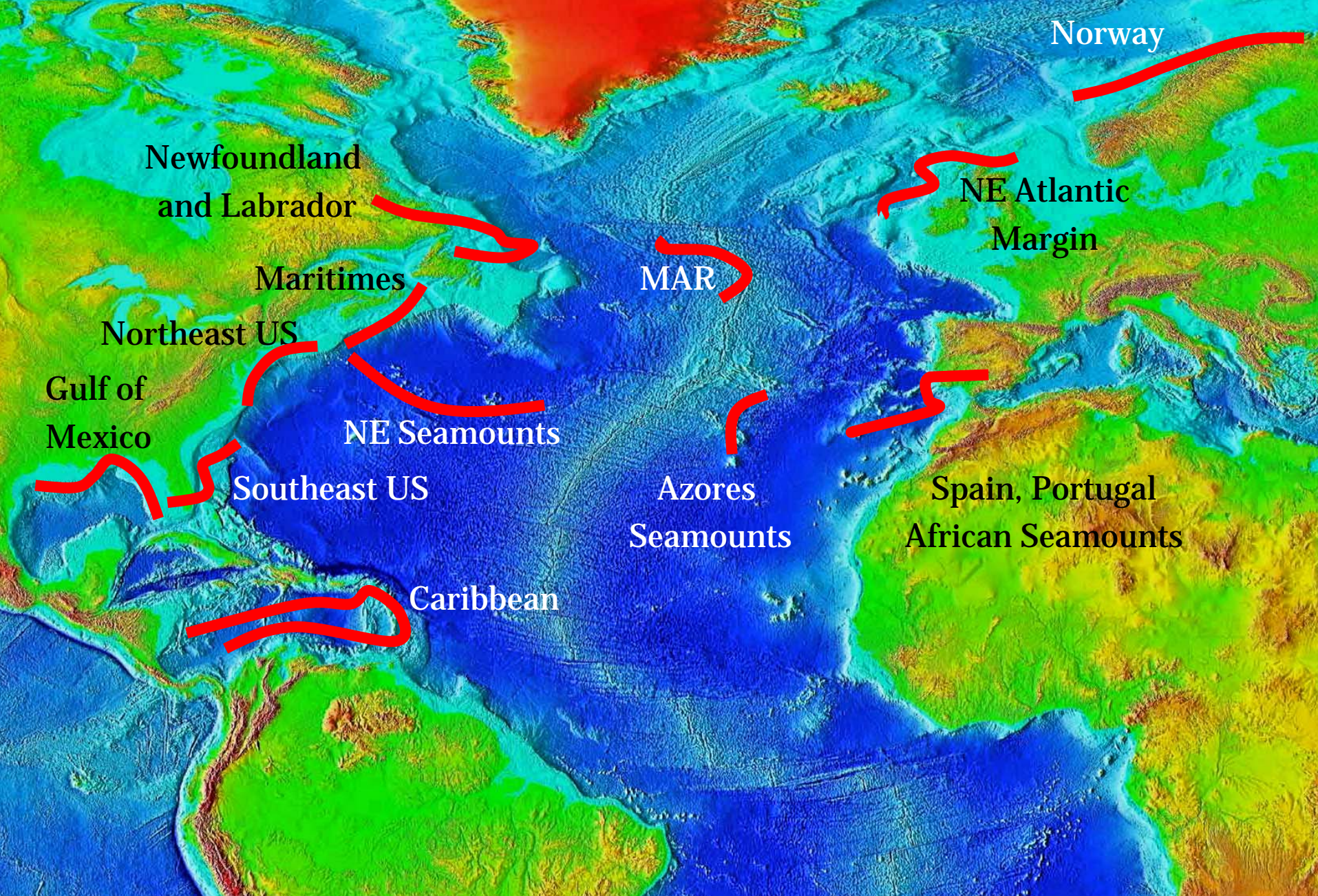
Population Connectivity-

“the dispersal, survival, and reproduction of migrants, so that they contribute to the local gene pool”

-Cowen et al., 2007

1. Linkages
2. Approaches
3. Challenges

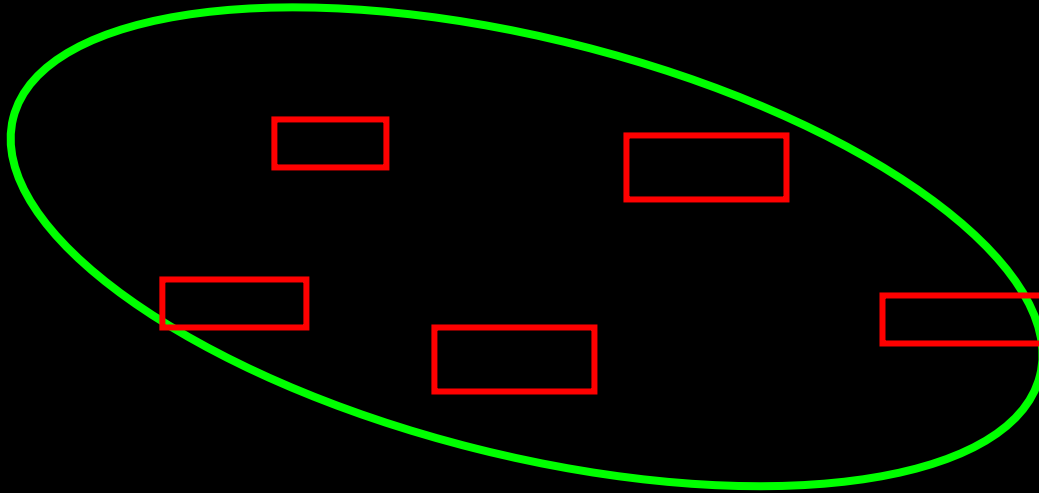




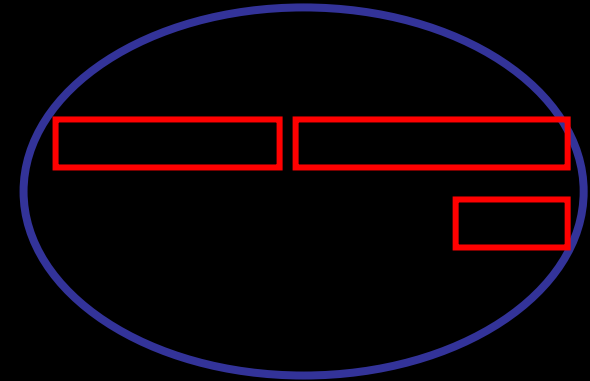
Cold water coral locations in the North Atlantic

North Atlantic Seamounts

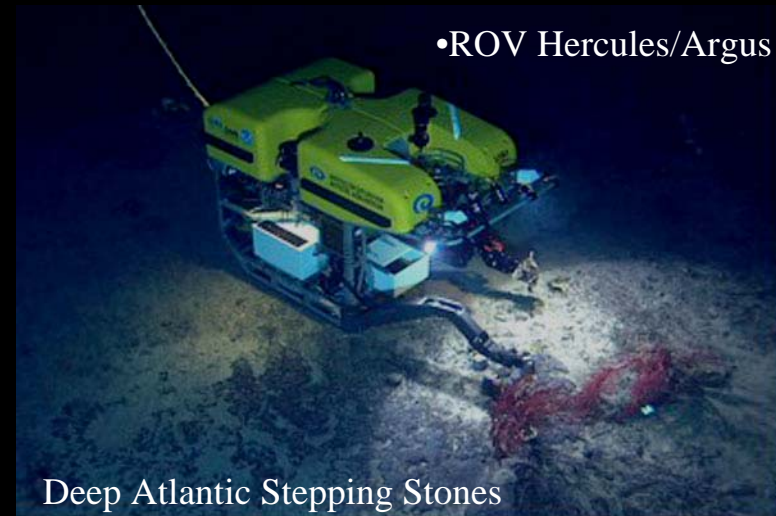
New England Seamount Chain



Corner Rise Seamounts



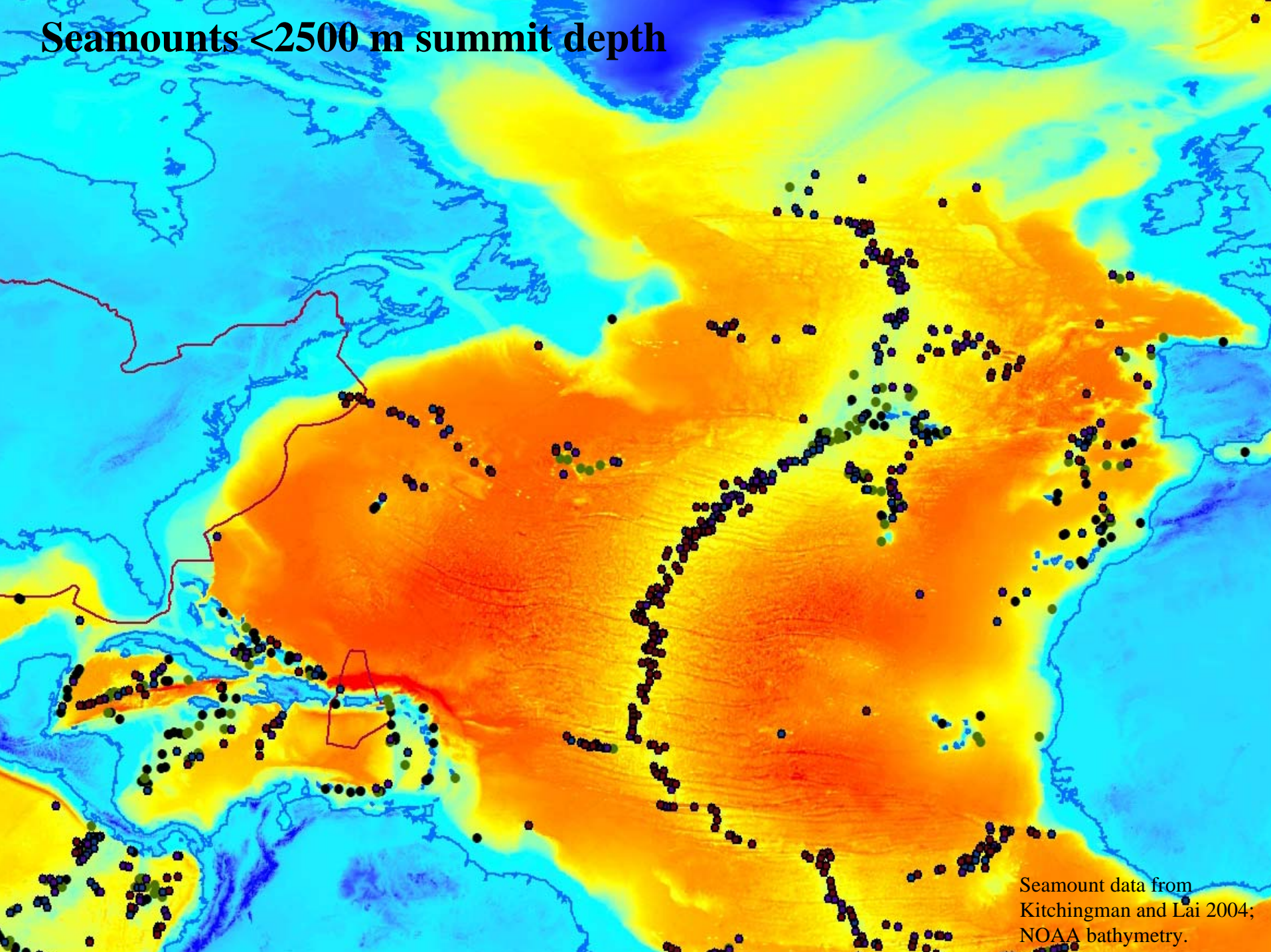
•ROV Hercules/Argus



Quick Survey Summary of the New England and Corner Rise Seamounts

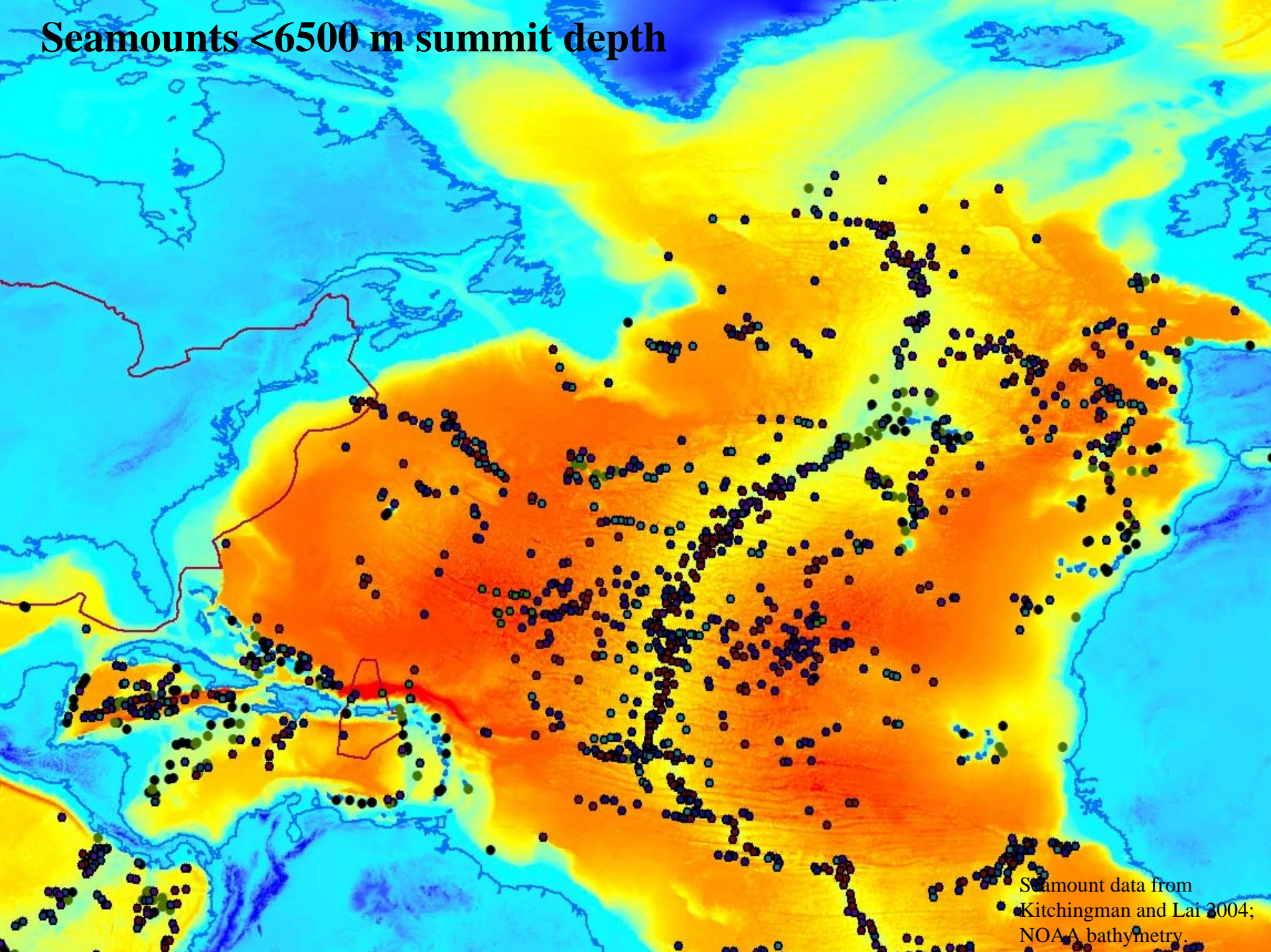
- Total of 61,714 observations
- 270 species, dominated by sponges, corals, echinoderms
- 60 species found only on the New England Smts
- 75 species found only on the Corner Rise Smts
- 135 species found at both the New England Smts & the Corner Rise Smts
- Host associate relationships- “facultative to obligate”

Seamounts <2500 m summit depth

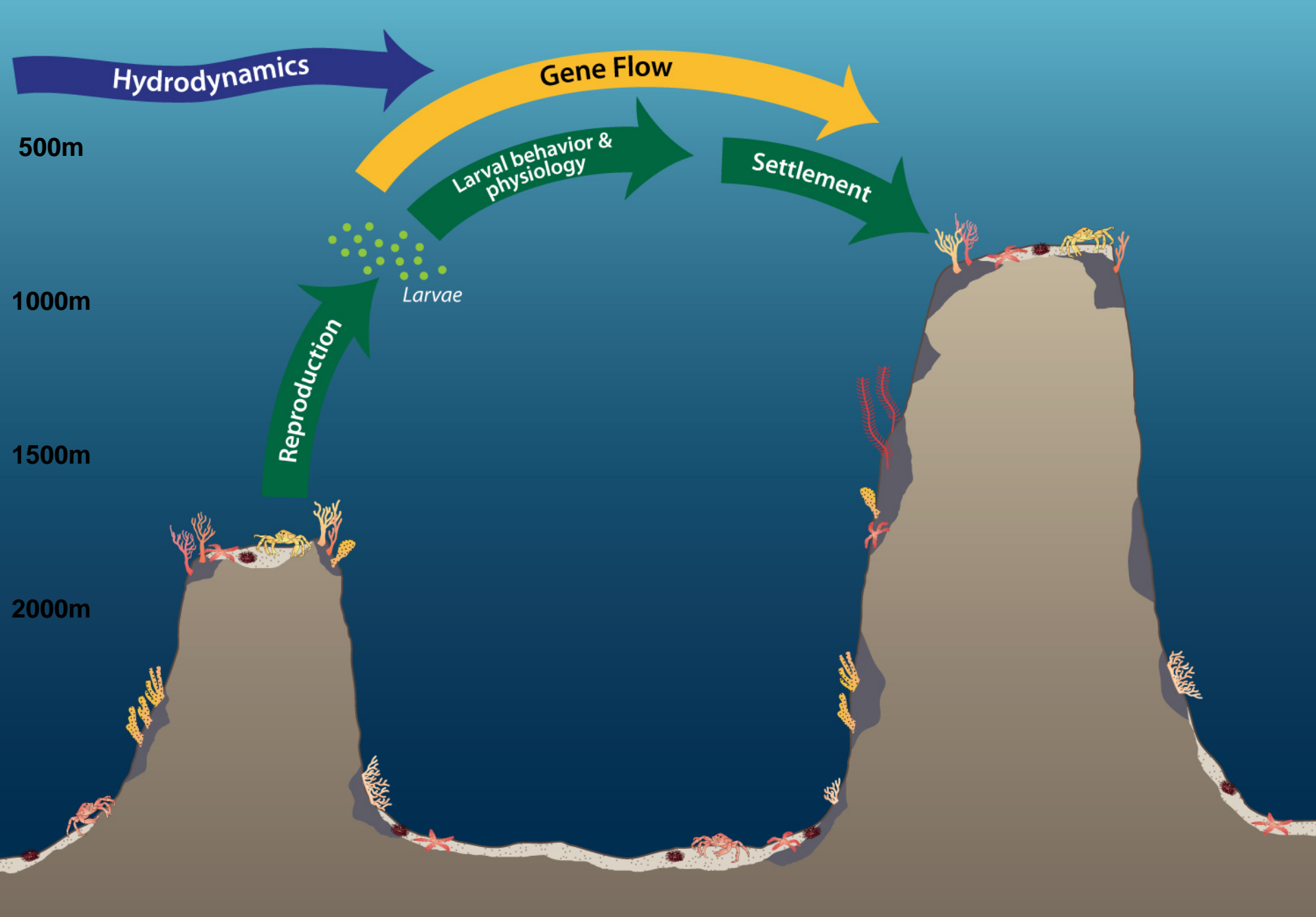


Seamount data from
Kitchingman and Lai 2004;
NOAA bathymetry.

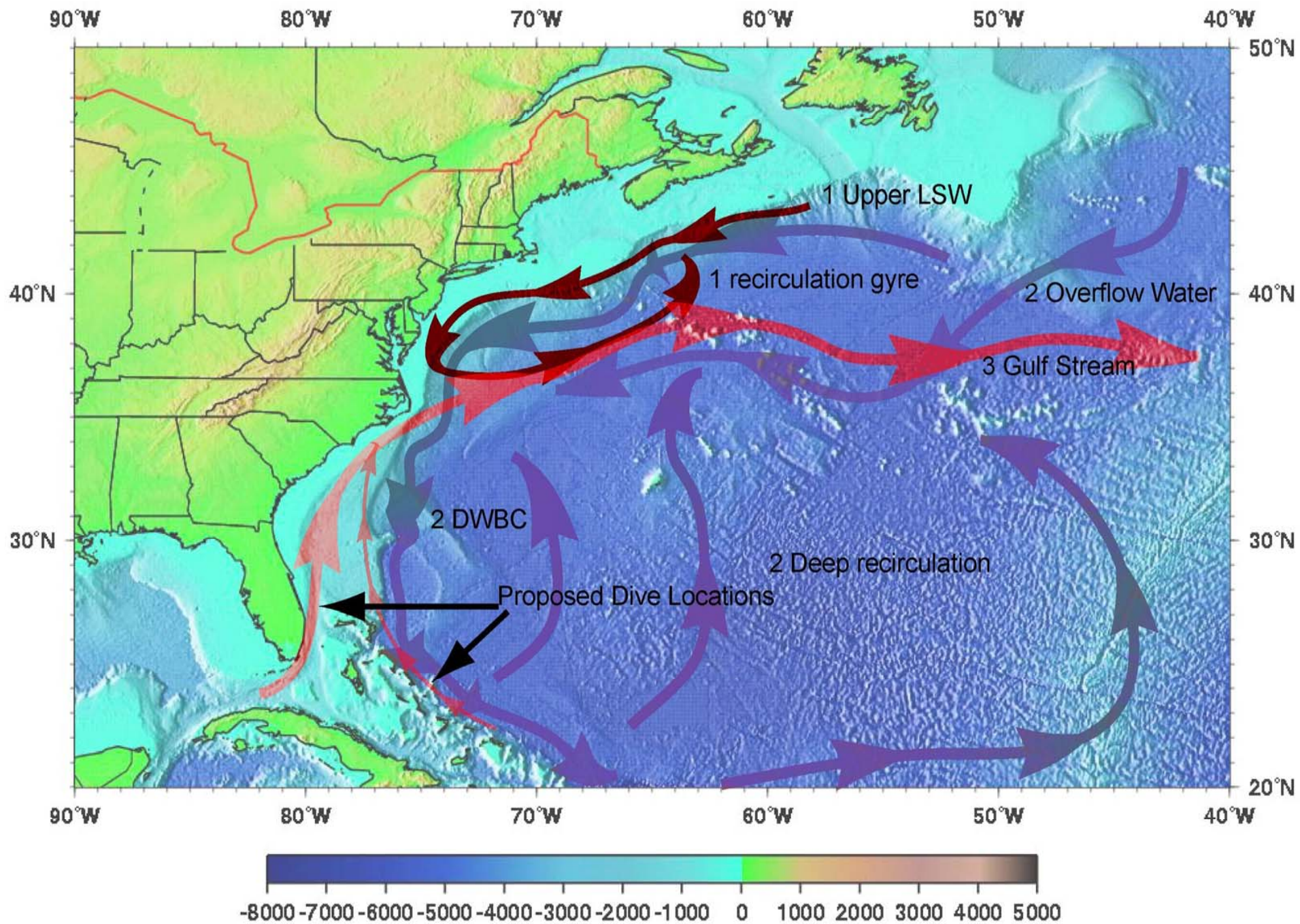
Seamounts <6500 m summit depth

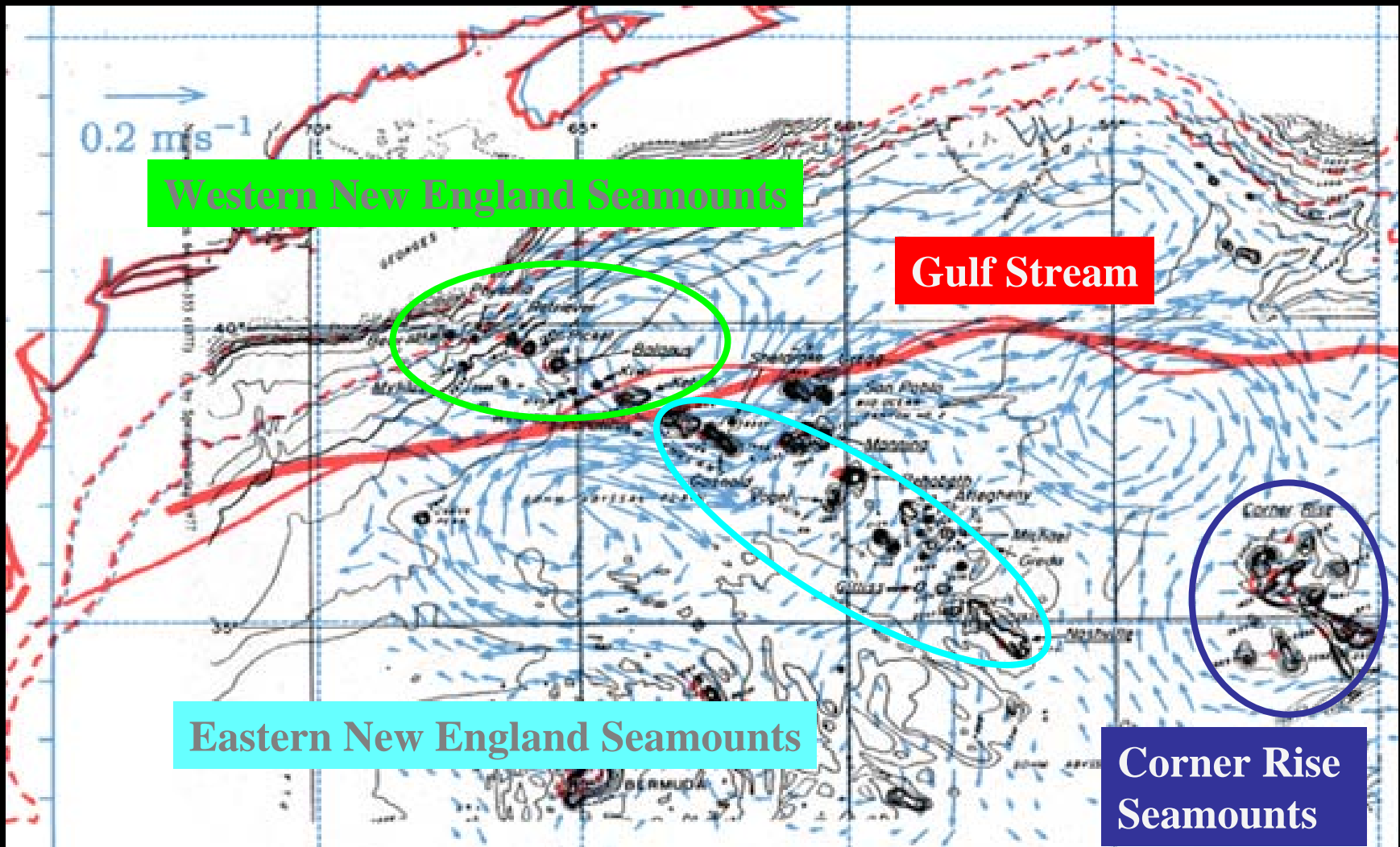


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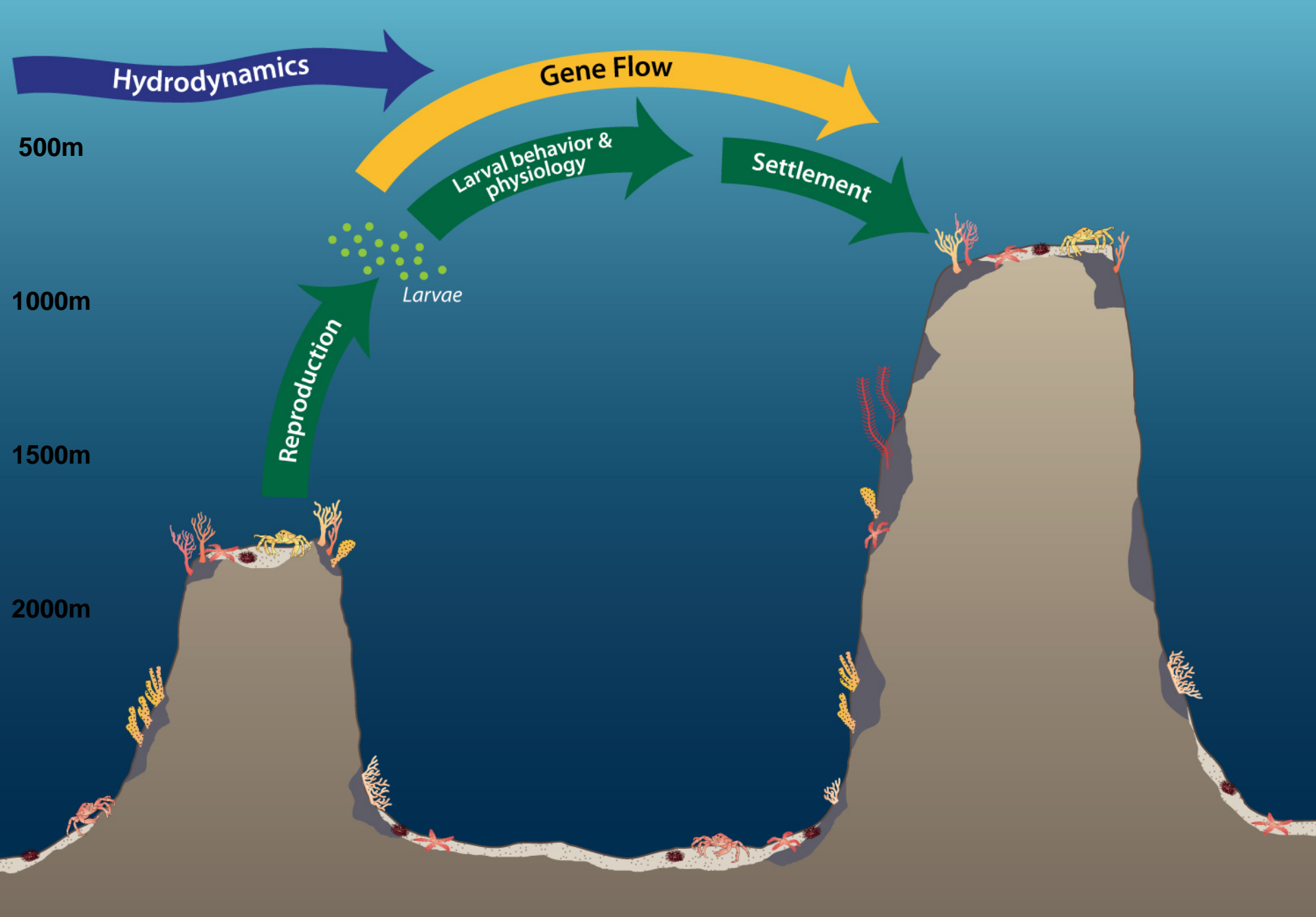
Seafloor morphology, Habitat availability(abiotic/biotic), Depth, Reproduction/Settlement, Hydrodynamics



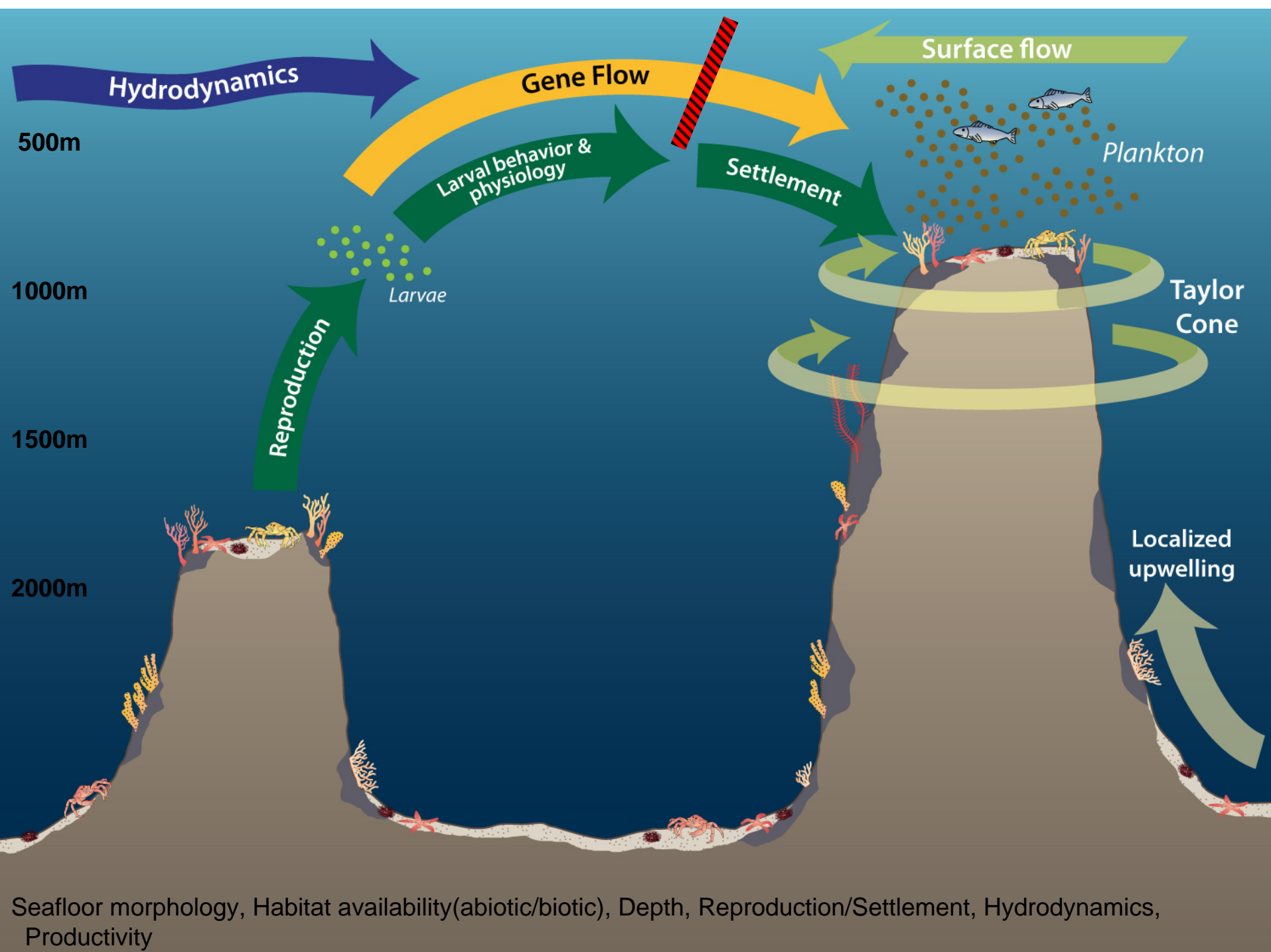


Predicted flow (blue arrows) of the Gulf Stream at 3000m depth (mean flow in solid red) across the New England Seamounts

(calculated from surface to deep water models; modified from Houghton et al., 1977; Qiu, 1994)



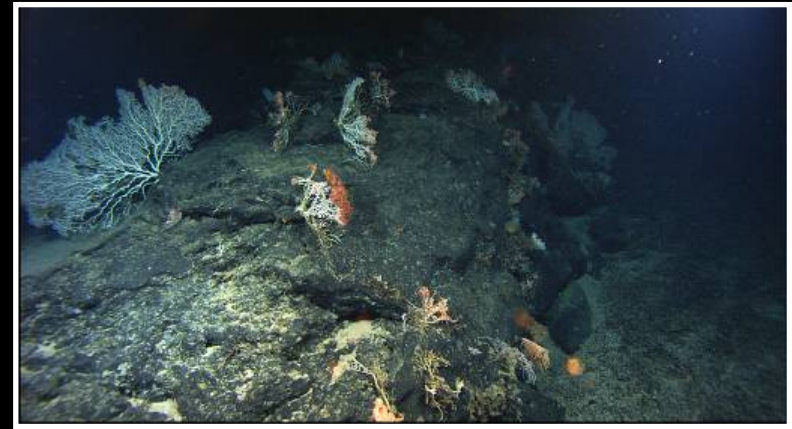
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Population Connectivity-

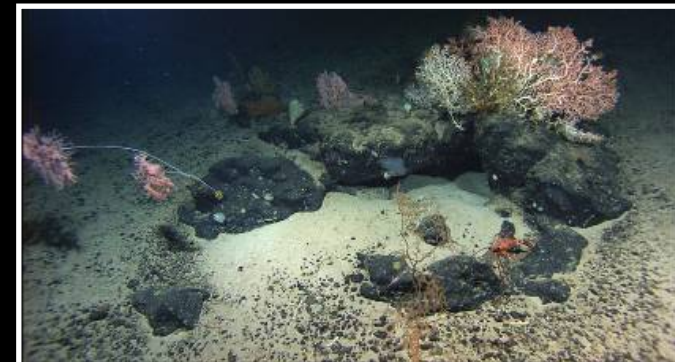
“the dispersal, survival, and reproduction of migrants, so that they contribute to the local gene pool”

-Cowen et al., 2007



Basic approaches to inferring connectivity:

- 1) indirect (inferring effective migration)
 - Genetic methods to estimate gene flow from genetic differences among populations (Slatkin et al., 2003; Benzie 1999); different gene regions evolve at different rates so must pick markers yield appropriate scale^L
- 2) direct (assigning sources populations, natal origins, or parents)
 - Larval and adult tags (Jones et al., 1999; Alamay et al., 2007)
 - Fish otolith chemistry (Thorrold et al., 2001)
 - Genetic markers for assignment of individuals to populations
(Mantel et al., 2005; Jones et al., 2005)





First things first:

Must identify what a species is...

- DNA sequences reveal cryptic species
 - Species morphologically indistinguishable
 - DNA data assists with taxonomy
 - France (2007): example in bamboo corals
- DNA sequence data
 - octocorals: *msh1* gene*
 - black corals: mitochondrial intergenic*
- sequenced the complete mitochondrial genomes of 2 bamboo corals and discovered novel 5-gene inversion
 - Brugler & France (in review)

* The standard mtCOI barcode is not sufficiently variable in corals.



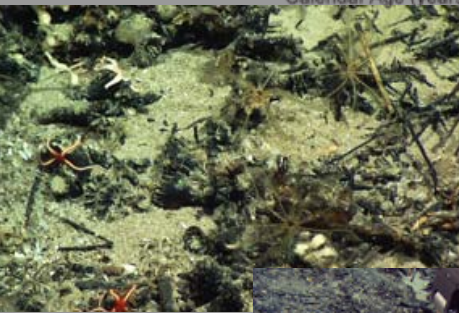
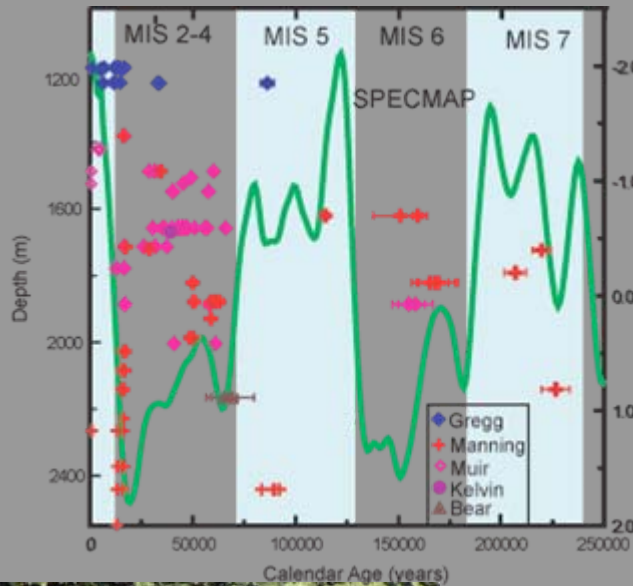
Phylogeographic analyses of abundant taxa

- *Paramuricea* spp. (octocoral)
- *Metallogorgia melanotrichos* (octocoral)
- *Parantipathes* spp. (black coral)

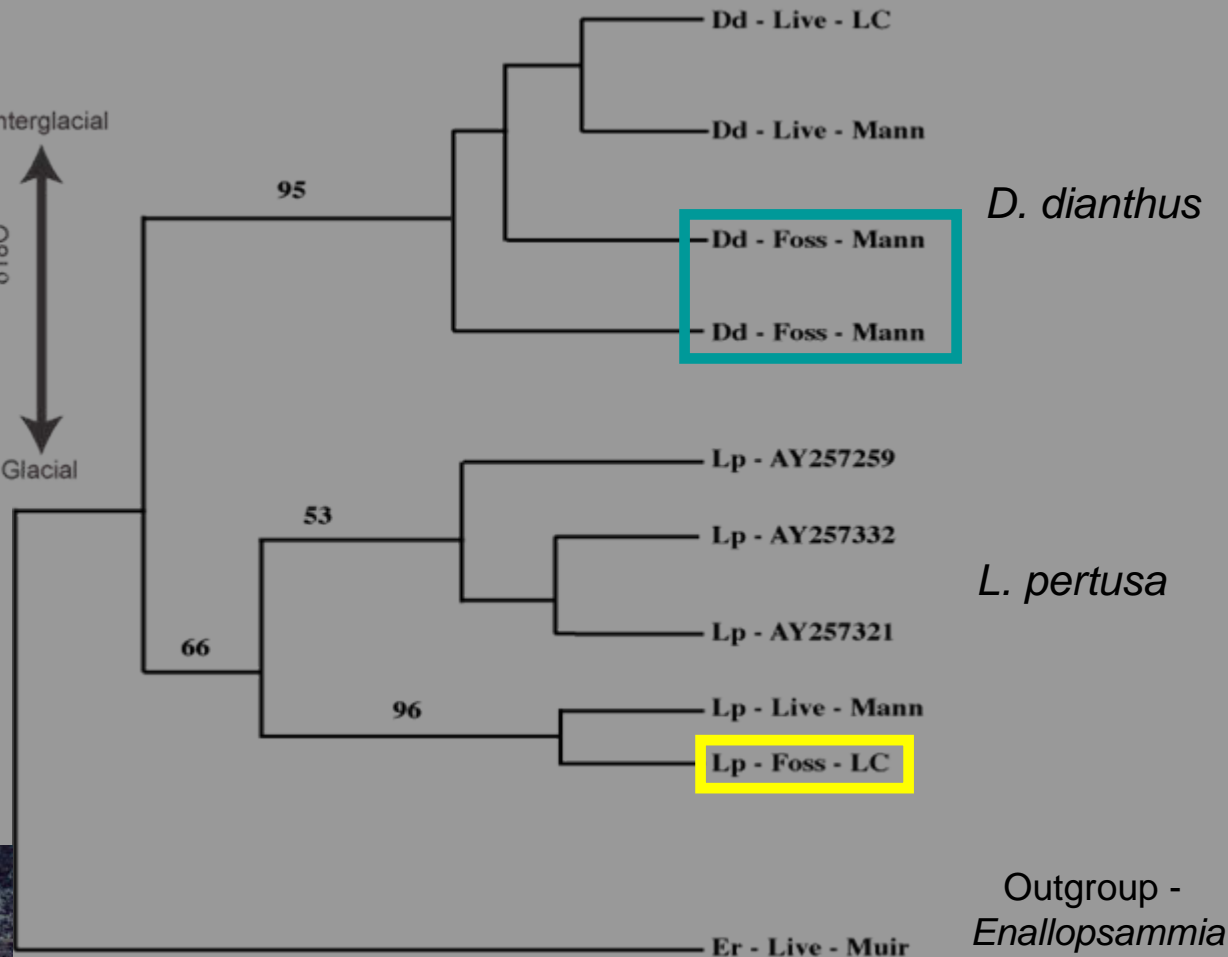
***Paramuricea* sp.**

Goode Peak, Corner Seamount, 1916 m

Paleo-Population Genetics - Connectivity and Climate Change



Fossil *D. dianthus*



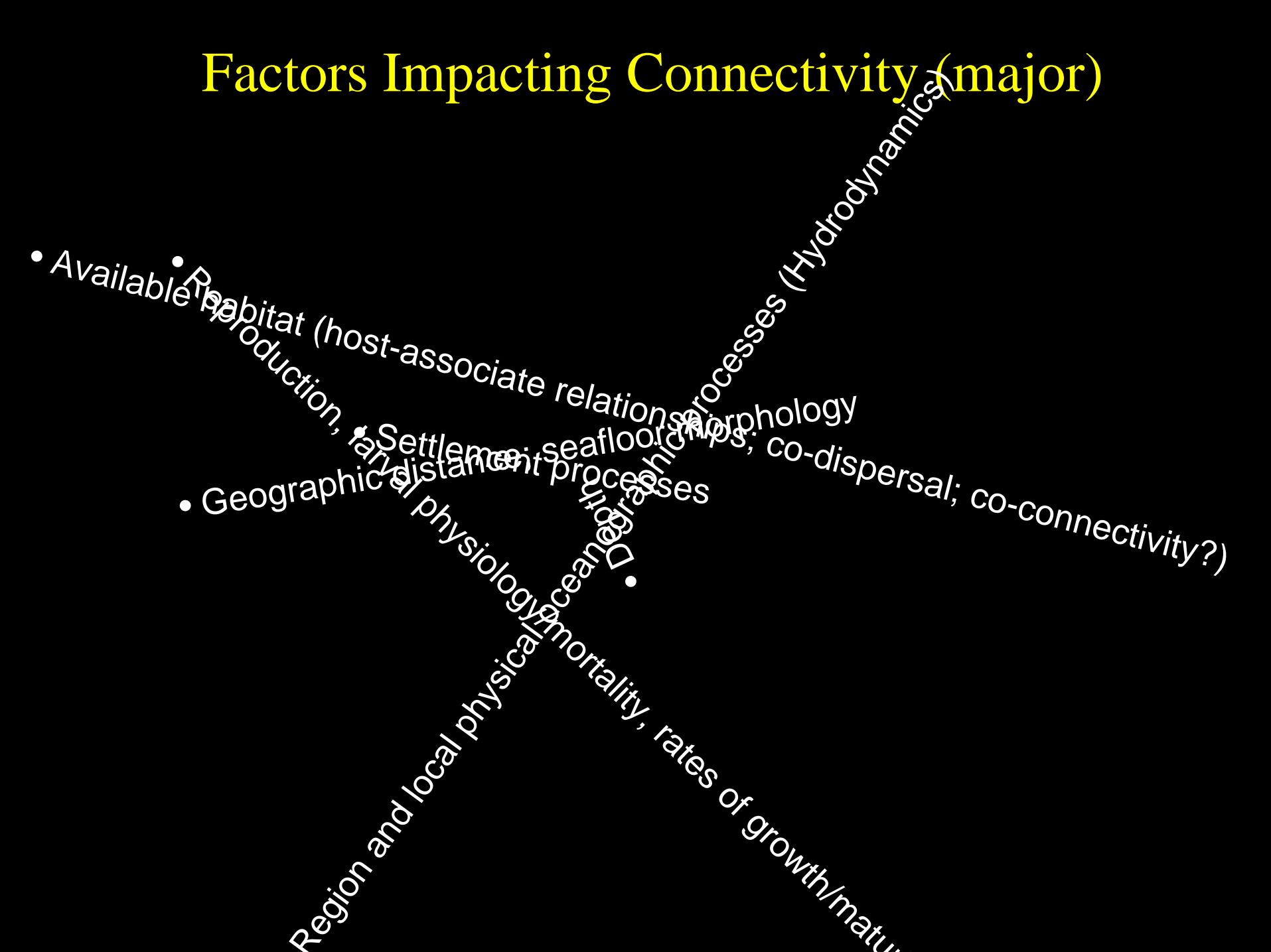
- Preliminary Results -
- ITS1, ITS2 compared to live sequences (~200bp)



Factors Impacting Connectivity (major)

- Geographic distance; seafloor morphology
- Available habitat (host-associate relationships; co-dispersal; co-connectivity?)
- Depth
- Region and local physical oceanographic processes (Hydrodynamics)
- Reproduction, larval physiology/mortality, rates of growth/maturity
- Settlement processes

Factors Impacting Connectivity (major)



Challenges for understanding “Connectivity”

1. Identifying the dynamic temporal and spatial scales over which all of these processes interact
2. Designing the appropriate sampling design to address questions of connectivity on these scales
3. Identifying species through morph. & molecular taxonomy
4. Identifying population/phylogeographic boundaries
5. Obtaining enough samples/individuals for population genetics
6. Finding the appropriate genetic marker for the question
7. Obtaining co-located/co-incident physical and environmental data

Major Research Questions

(Linkages and Connectivity)

- 1. What are the dominant processes that structure differences in connectivity in Atlantic coral ecosystems?
- 2. What are the key temporal and spatial scales over which these processes operate...to cause differences in connectivity in coral ecosystems?
- 3. How do we overcome the key technical (e.g., genetic resolution, sampling designs) problems assessing connectivity and gene flow (over spatial and temporal scales)?
- 4. What are the impacts of fisheries on altering coral ecosystem connectivity?

What Do We Need?

- to realize we can not extrapolate current dispersal/connectivity results of any taxa or seamount to another. For example, from the South Pacific or the Equatorial Pacific
- to realize that species characterized as having “limited larval dispersal” can travel large oceanic distances
- to realize that isolated coral communities are only isolated if larval dispersal and colonization limit the exchange of genetic material (distance \neq isolation)
- to realize that any plan to assess coral community diversity, endemism, and connectivity “must” include the integrated use of genetic approaches