

The University of Maine

DigitalCommons@UMaine

Annual Maine Aquaculture R&D and Education
Summits

Conferences and Summits

Winter 1-17-2020

The Maine e-DNA EPSCoR Program and its Ties to Maine Salmon Ecosystems

Michael T. Kinnison

David Emerson

Heather Leslie

Kate Beard-Tisdale

Kody Varahramyan

Follow this and additional works at: https://digitalcommons.library.umaine.edu/ari_rd-ed

This Presentation is brought to you for free and open access by DigitalCommons@UMaine. It has been accepted for inclusion in Annual Maine Aquaculture R&D and Education Summits by an authorized administrator of DigitalCommons@UMaine. For more information, please contact um.library.technical.services@maine.edu.

The Maine-eDNA EPSCoR Program and its Ties to Maine Salmon Ecosystems

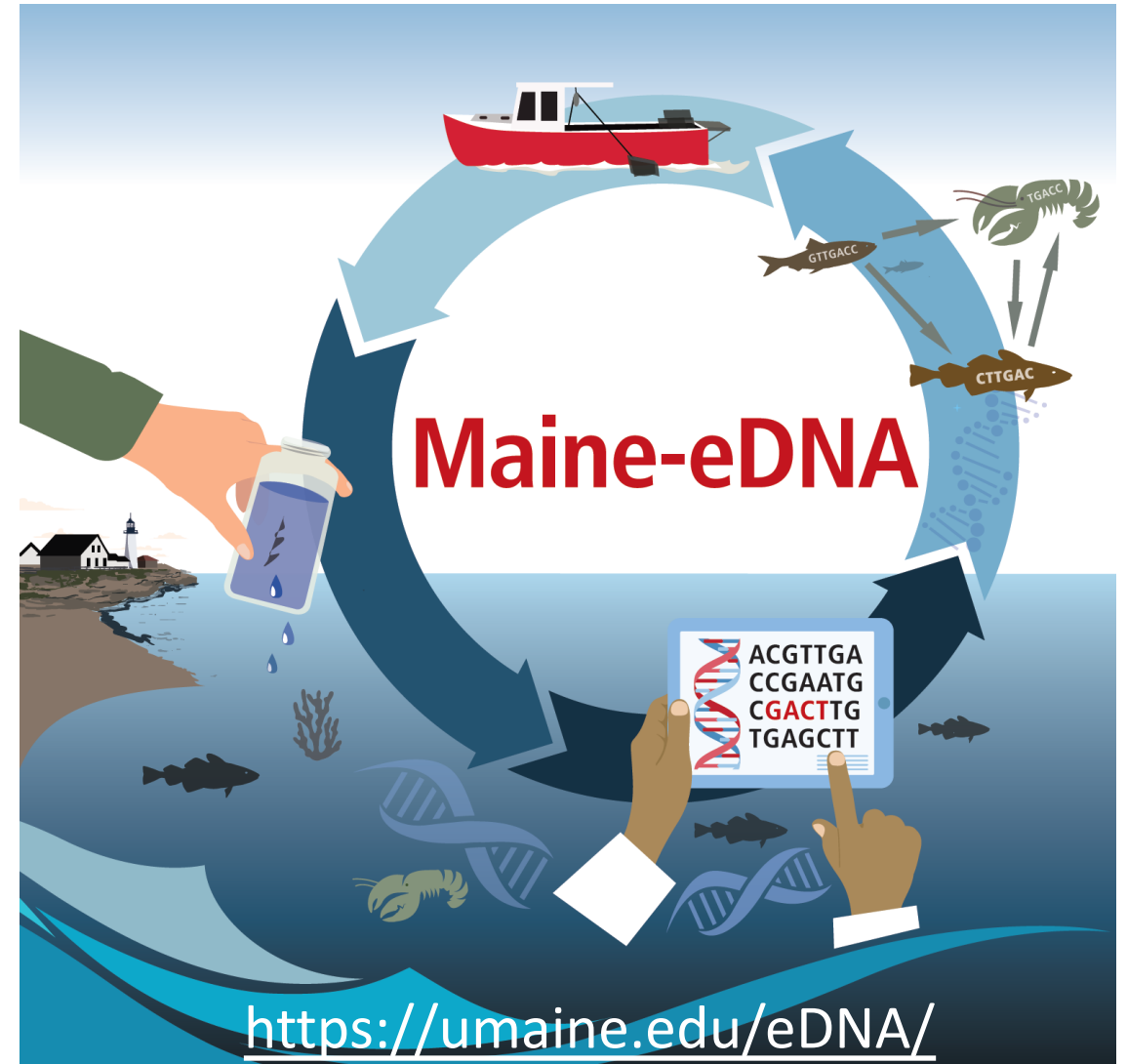
Michael Kinnison – UMaine SBE

David Emerson – Bigelow Laboratory

Heather Leslie – UMaine Darling Center

Kate Beard-Tisdale – UMaine CIS

Kody Varahramyan – UMaine VPR DGS



Environmental DNA (eDNA)

Environmental DNA (eDNA) – genetic material obtained directly from environmental samples (sediment, water, etc.) without any obvious signs of biological source material

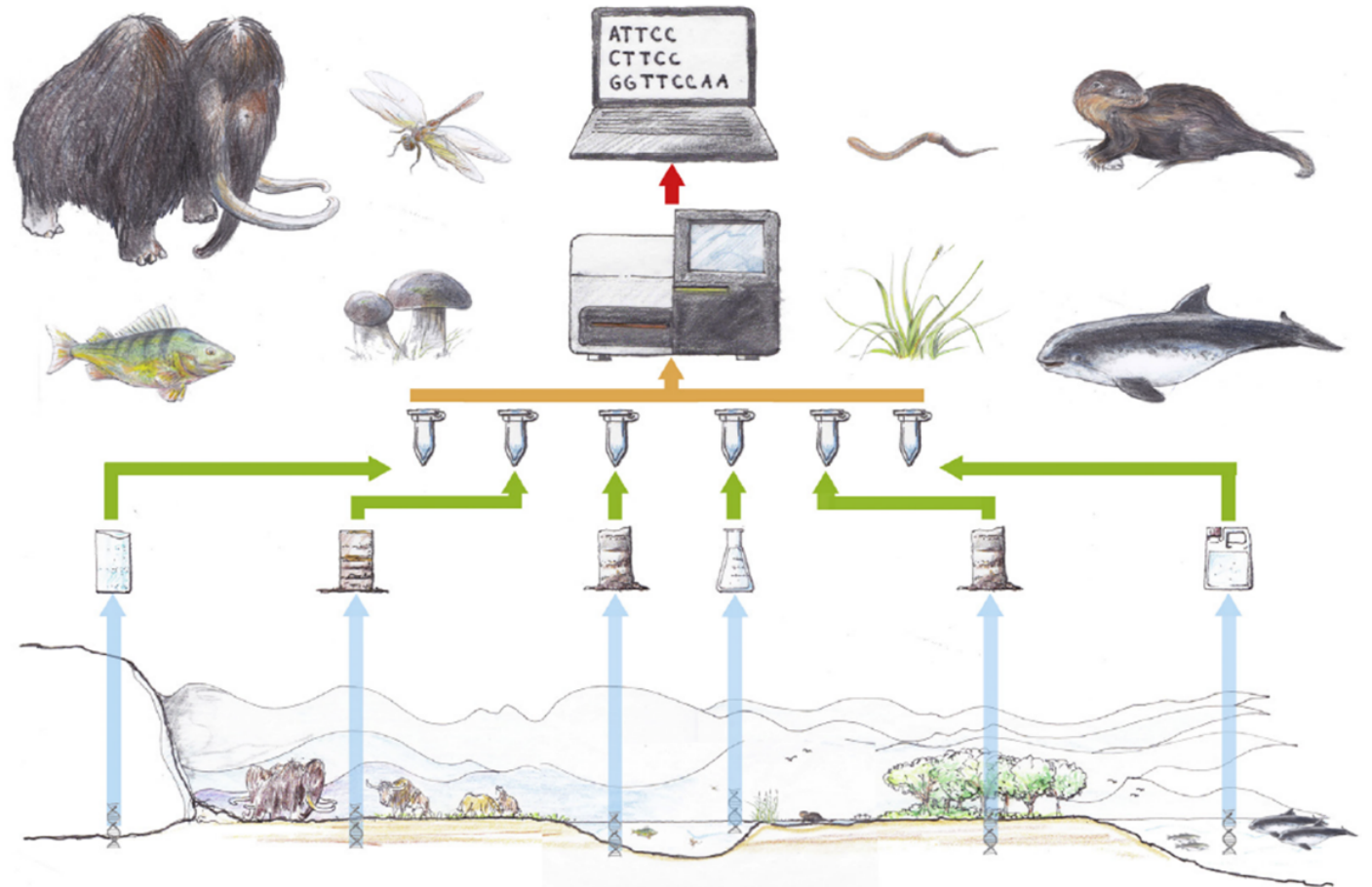
Thomsen and Wilerslev (2015), Biological Conservation 183, pp 4-18

Sources of eDNA:

- Cellular decomposition
- Whole shed cells
- Whole microorganisms (e.g., larvae)

Uses of eDNA:

- Targeted detection
- Targeted quantification
- Community characterization





eDNA Uses to Date

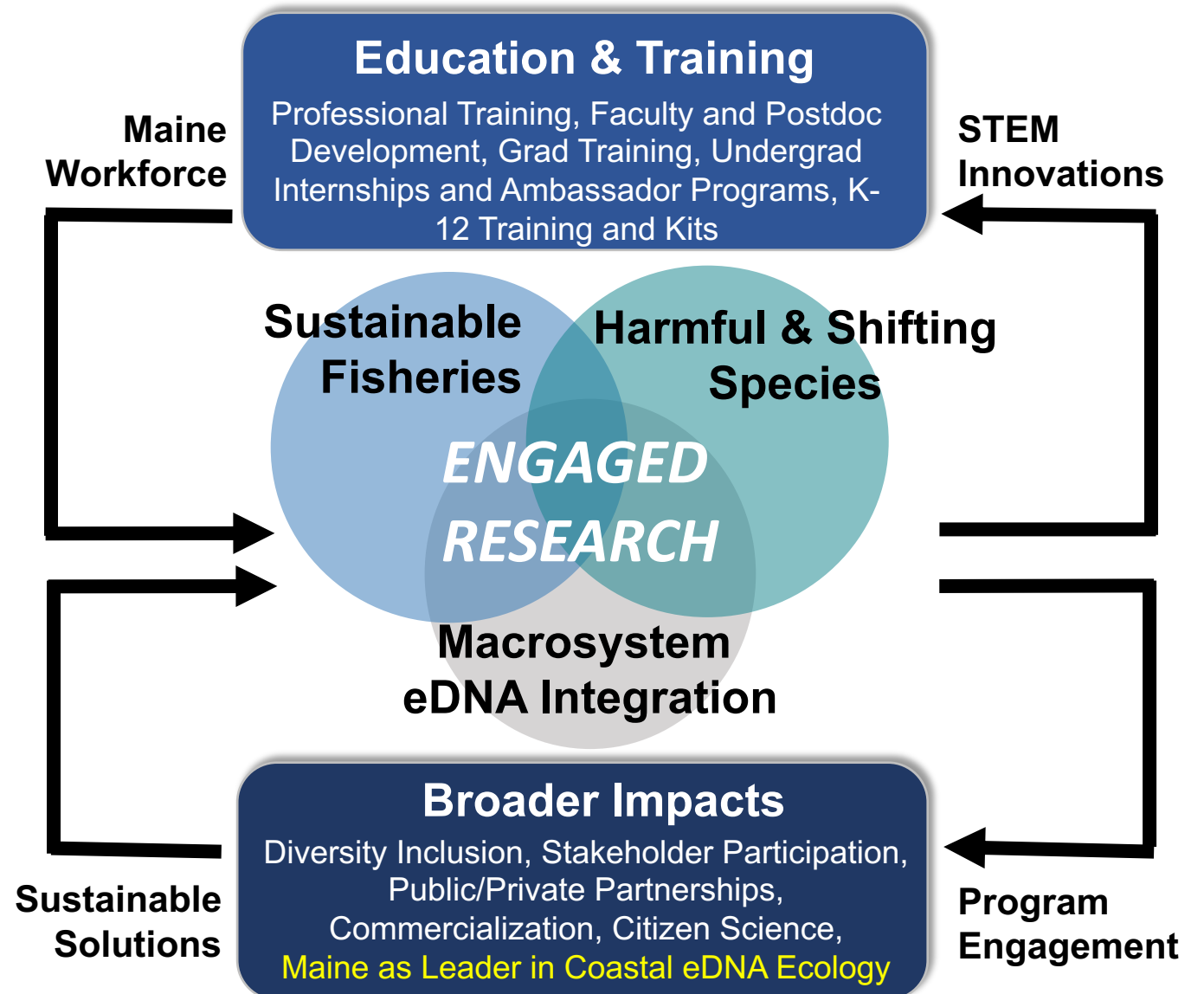
- ID species presence/absence, overall diversity
- Reveal distribution patterns over broad scales of time and space
- Detect rare, invasive, or unseen species
- Reconstruct past communities (via ancient/preserved DNA – it's very stable)
- Estimate biodiversity – for ecosystem management etc...

Bourlat et al., 2013 (Table 3) & Thomsen & Willerslev, 2015 (Table 1)

What is the Maine-eDNA EPSCoR Program?

**A \$20 million NSF EPSCoR
Research Infrastructure
Improvement Track-1 Award**

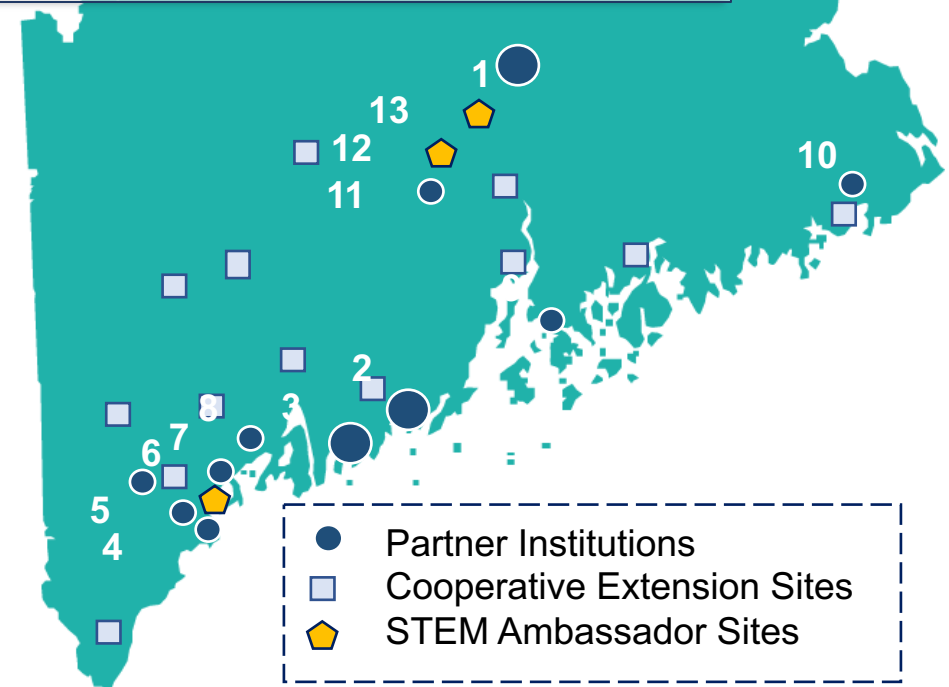
*“Molecule to Ecosystem:
Environmental DNA as a
Nexus of Coastal Ecosystem
Sustainability for Maine
(Maine-eDNA)”*



What is the Maine-eDNA EPSCoR Program?

- 32 Senior Personnel
- 3 ½ New Faculty Hires
- 21 Graduate Students
- 5 Postdocs
- >300 of Undergrad Interns
- 4 Primarily Undergrad Institutions
- 2 Research Nonprofits
- 3 Community Colleges
- 1 Land & Sea Grant University

1. University of Maine
2. Darling Marine Center (UMaine)
3. Bigelow Laboratory
4. University of New England
5. Southern Maine Community College
6. Gulf of Maine Research Institute
7. University of Southern Maine
8. Maine Technology Institute
9. Maine Maritime Academy
10. University of Maine at Machias
11. Kennebec Valley Community College
12. Colby College
13. Eastern Maine Community College



What is the Maine-eDNA EPSCoR Program?

VISION

Make Maine *'the DNA Coast'* - a world leader in eDNA-based partnerships, understanding, and sustainability of coastal marine and freshwater ecosystems.

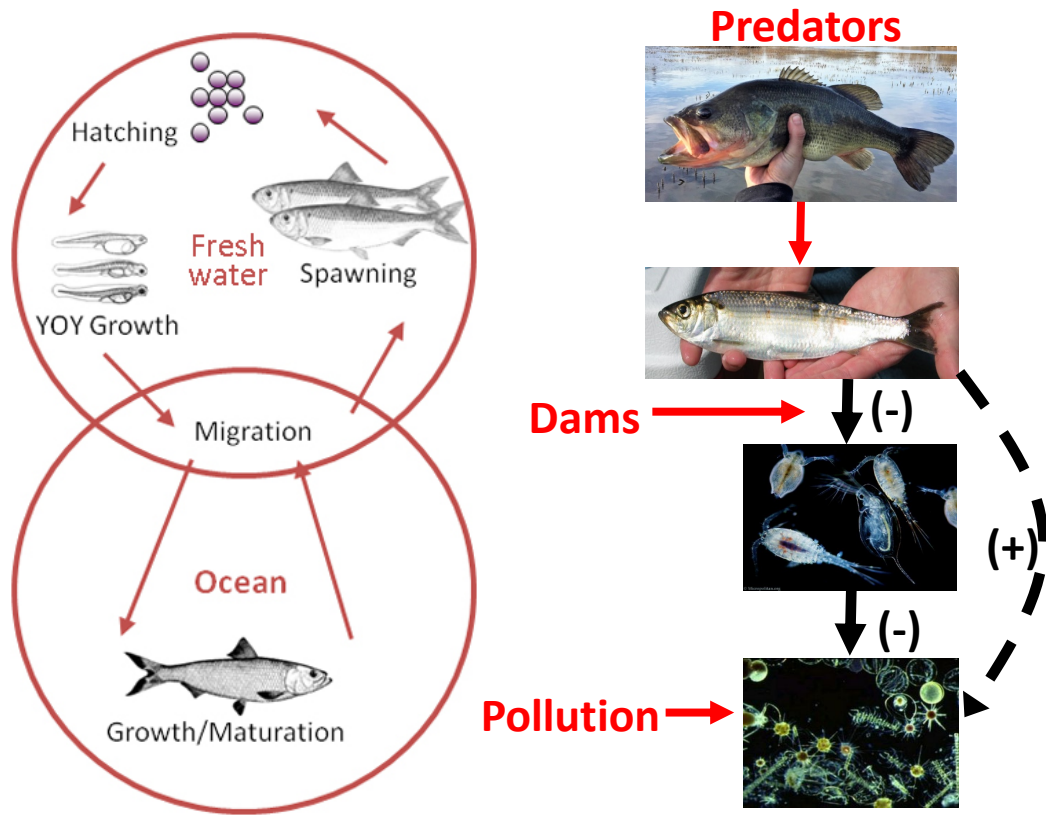
MISSION

Harness the power of eDNA science to:

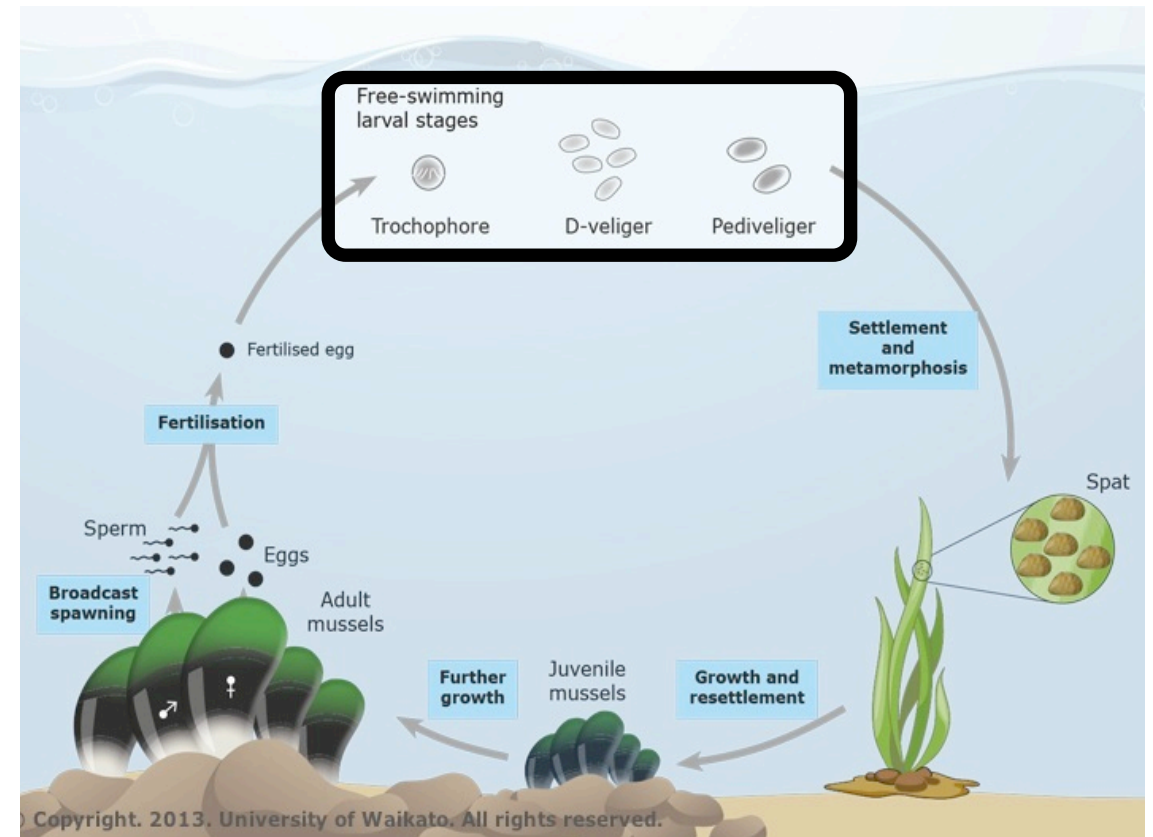
- 1) Advance ecological understanding crucial to the current needs of Maine's marine and freshwater resources.**
- 2) Build the Big Data and IP innovations, technical workforce, and partnership capacities to address increasingly large-scale and complex sustainability challenges of changing coastal ecosystems.**

Theme 1: Sustainable Fisheries

Goal 1.1: Ecosystem-Based Restoration: eDNA Past, Present, and Future



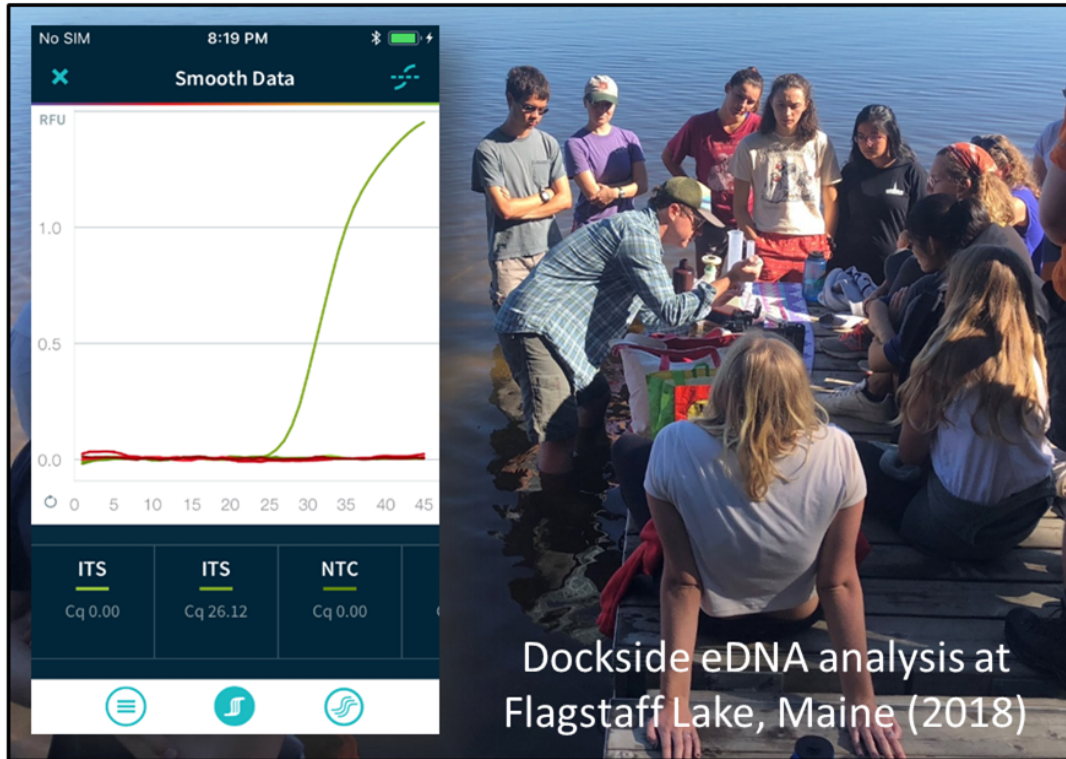
Goal 1.2: Larval Black Box: Revealing Early Life Ecology with eDNA



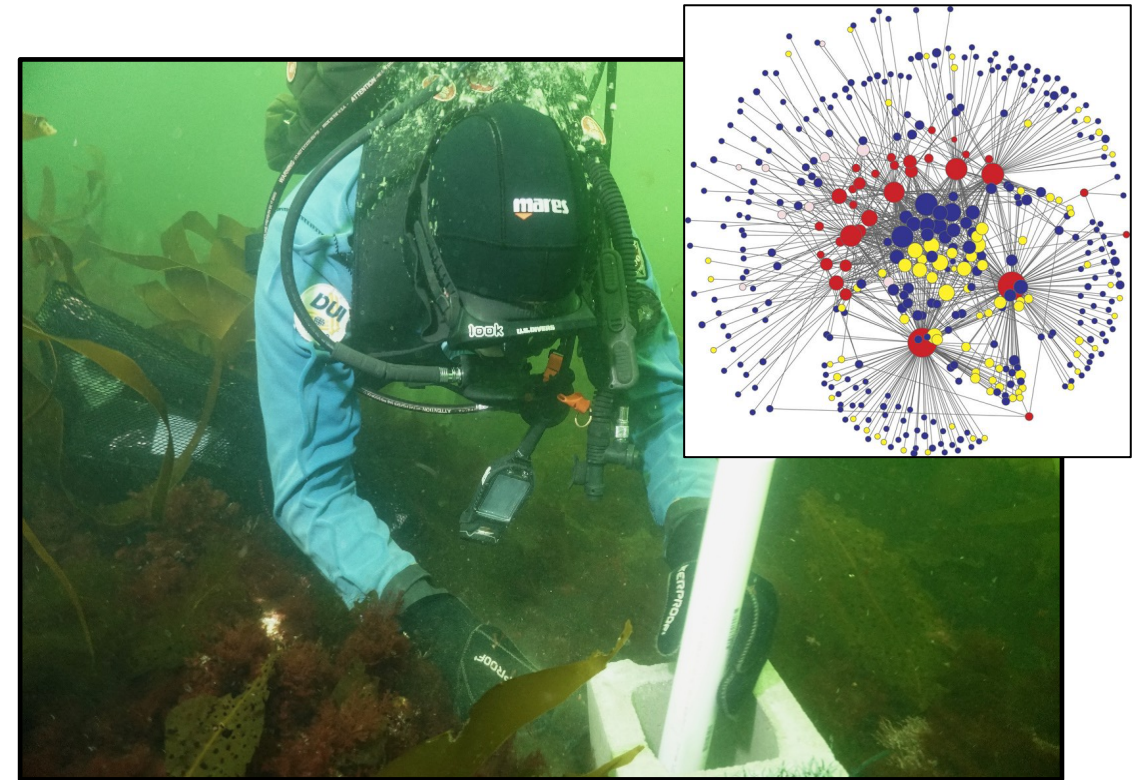
Integration Objective: eDNA framework for ecosystem-based fisheries and aquaculture.

Theme 2: Harmful & Shifting Species

Goal 2.1: Harmful Blooms: Population and Community Ecology of Toxic Threats



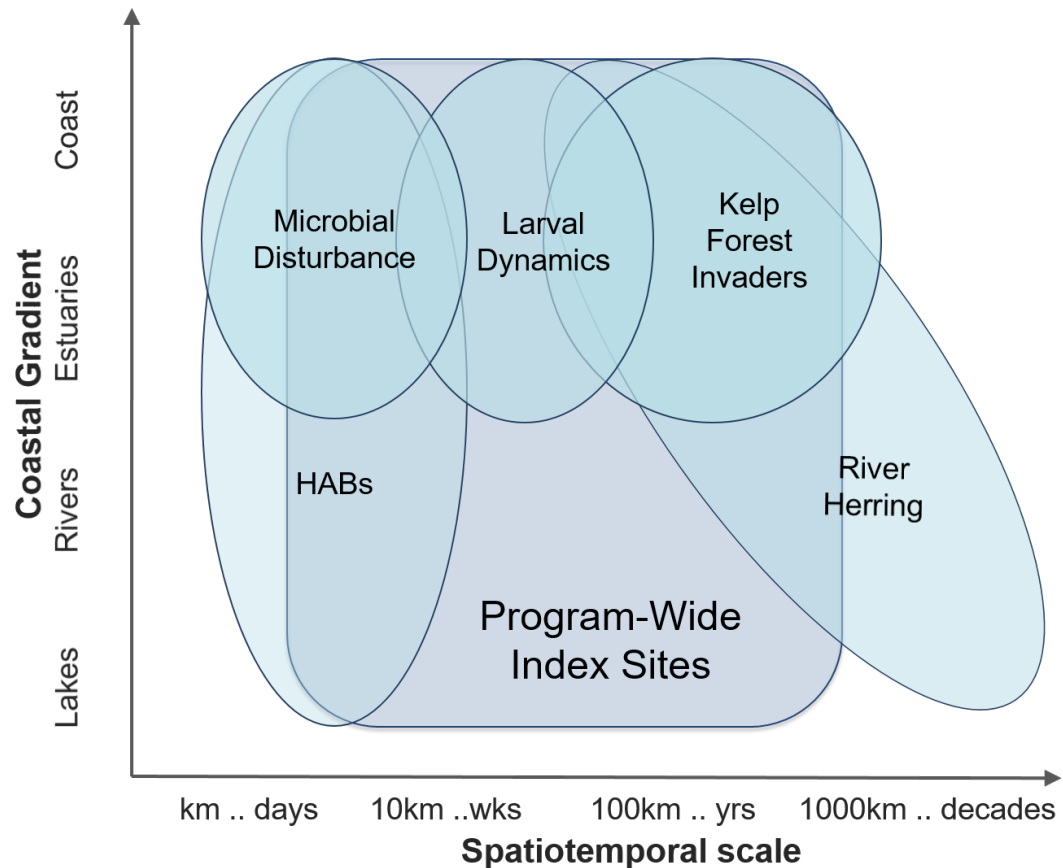
Goal 2.2: Species on the Move: Patterns and Determinants of Range Shifts



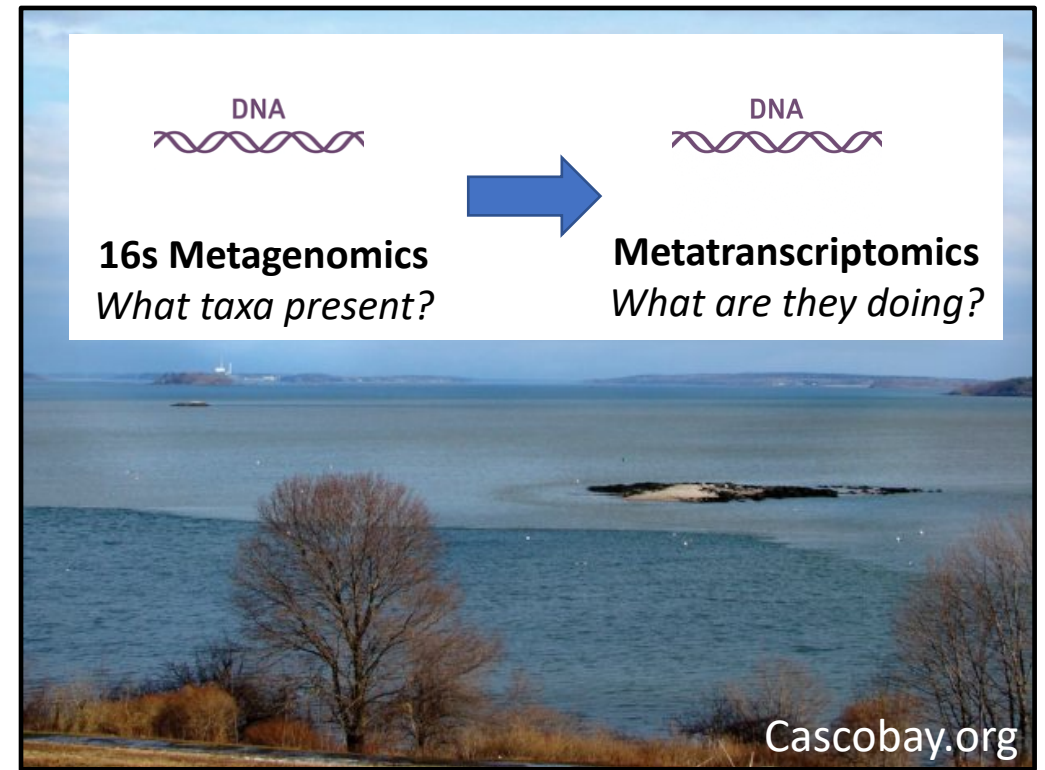
Integration Objective: eDNA early warning and forecasting tools to mitigate harmful effects

Theme 3: Macrosystem eDNA Integration

Goal 3.1: Big Data Integration: Dynamics and Stability of Maine's Coastal Macrosystem



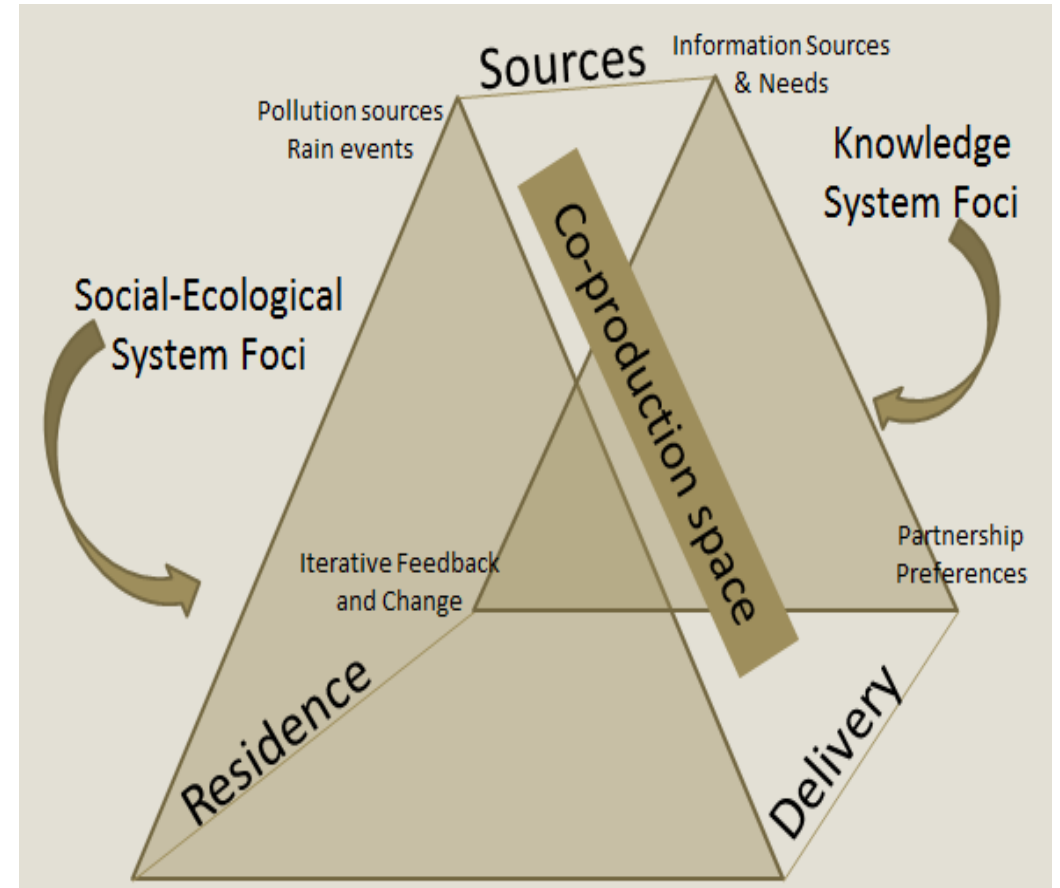
Goal 3.2: Microbial communities as biosensors of change



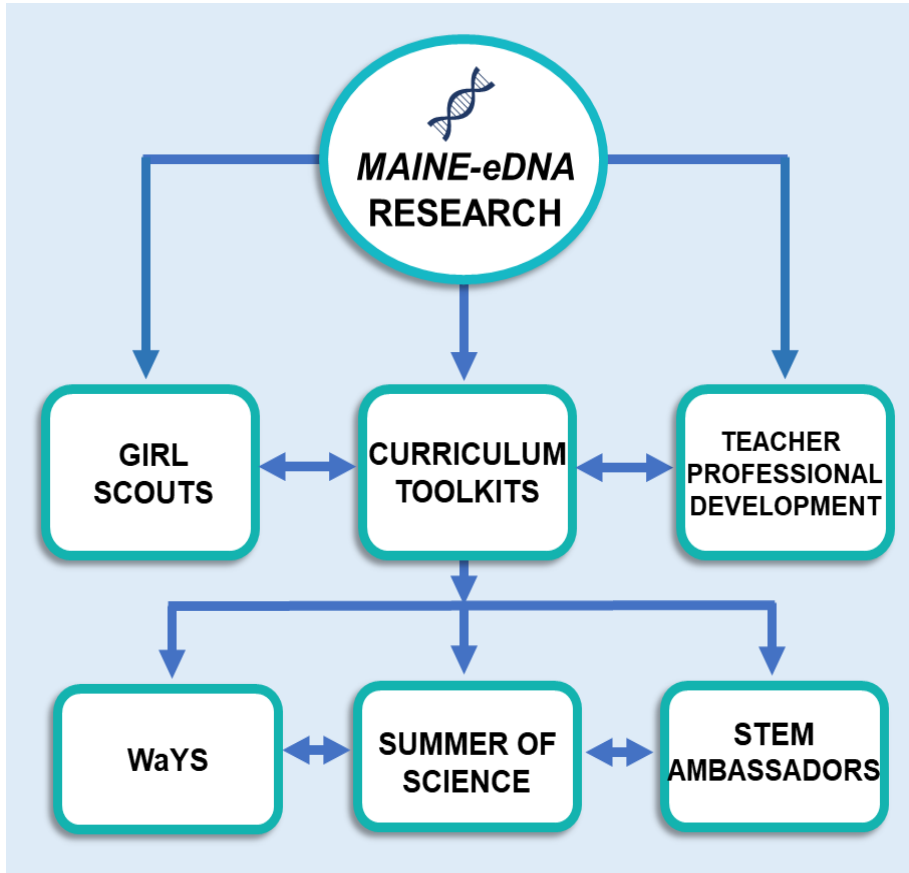
Casco Bay in 2014 after 5.6 inch rainfall

Theme 3: Macrosystem eDNA Integration

Goal 3.3 Communication Research: Comparative Study of eDNA Team Science



Education, Diversity and Workforce Development



K-PHD STEM Training



Environmental Professional Training



Citizen Science and Community Engagement

Aquaculture Relevant eDNA Applications

1. Developing eDNA tools to predict when/where 'seed' sets for scallops and mussels are expected
2. Quick (near real-time) tool for monitoring biosecurity on aquaculture farms
3. Improved tracking and monitoring of harmful algal bloom dispersal for informed farm ops decisions
4. Site selection to reduce exposure to pathogens, nuisance species, and predators with one tool



1. Larval Black Box

Relate scallop, *kelp*, *lobster*, and mussel gamete/*spore*/larval densities to eDNA counts as a function of *natal source* broodstock abundances, *fecundity*, *phenology*, and oceanic conditions.



Seasonal surface and benthic eDNA analysis (qPCR & eRNA)

- Plankton counts
- flow cytometry & microscopy
- Gonadal indices & sporogenesis
- Settlement collectors

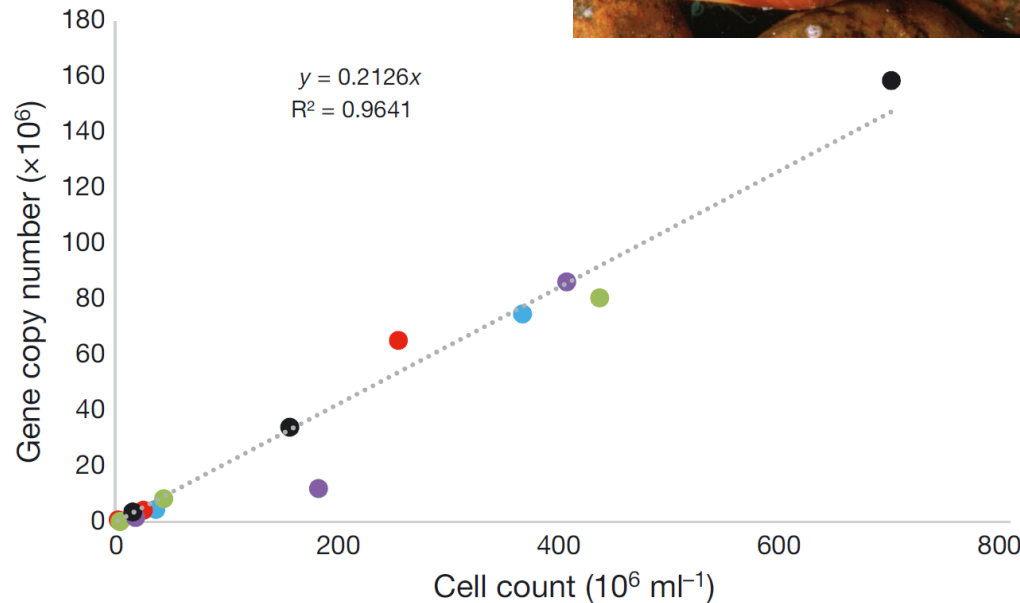


Fig. 2. Cell count of *P. magellanicus* sperm and resulting gene copy number from 5 dilution series (red, blue, green, purple and black filled circles)

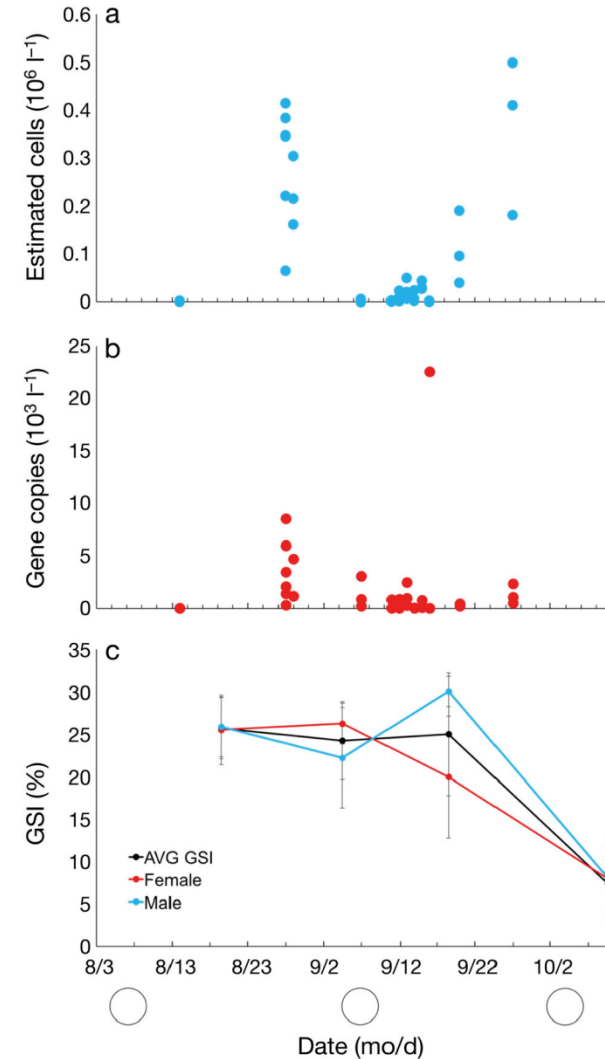
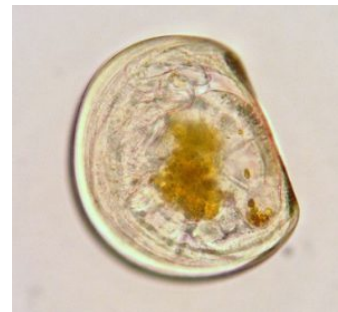
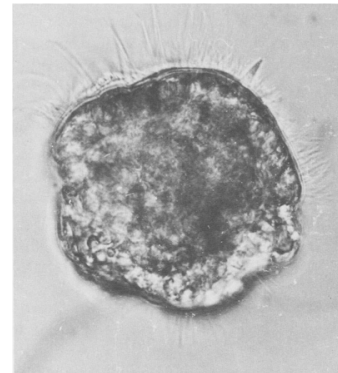
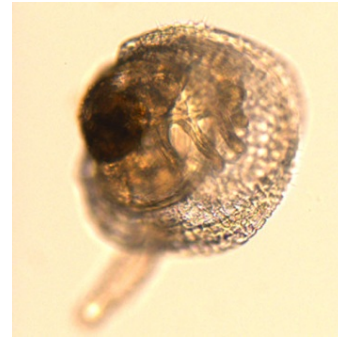
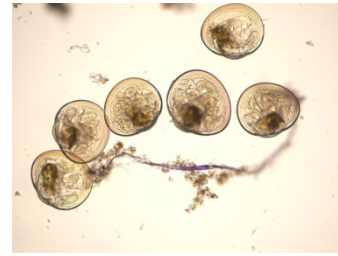
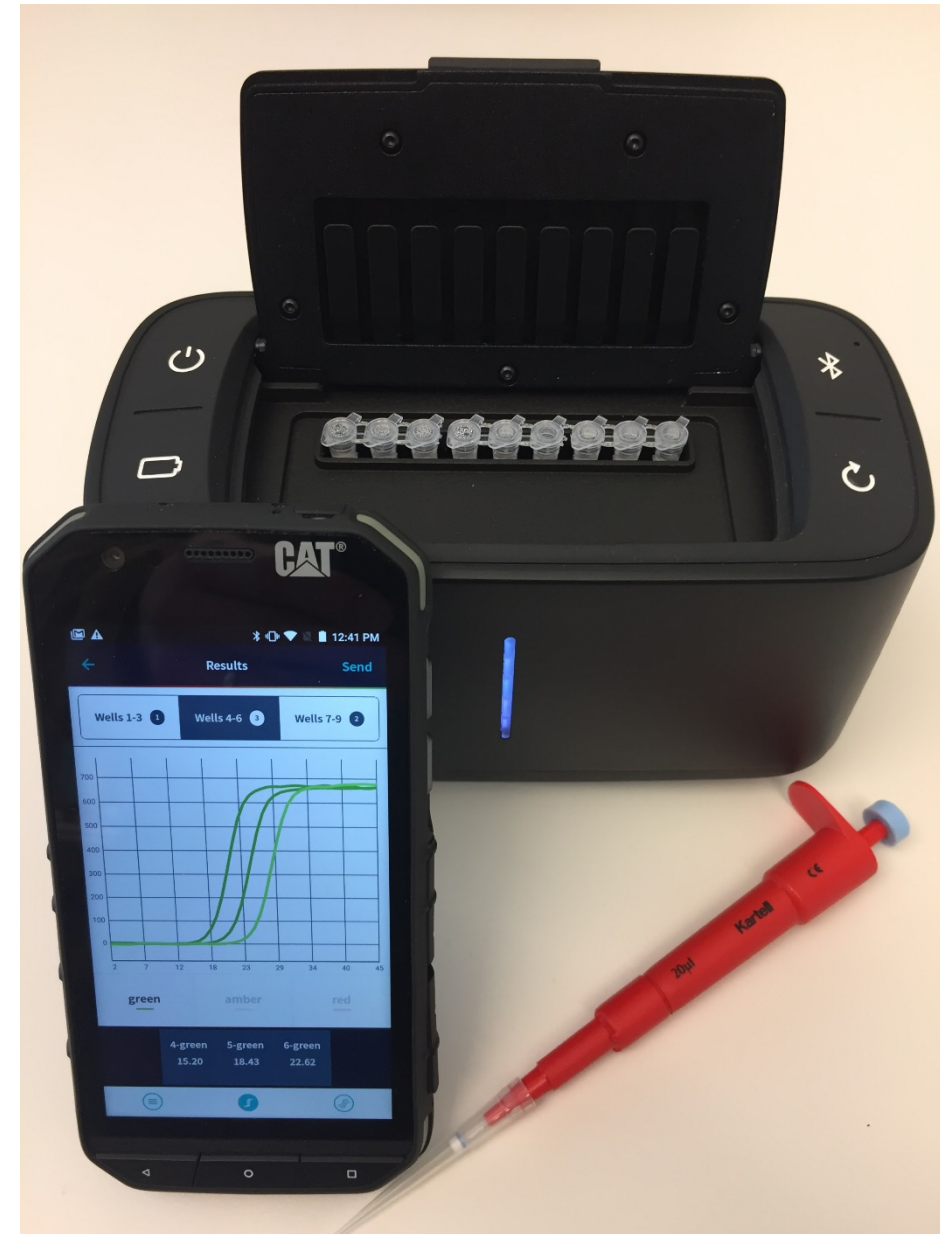


Fig. 3. Time series of (a) estimated sperm (cells < 20 μm) abundance and (b) gene copy number from cells > 20 μm from plankton pump samples collected < 0.5 m from dock-hung scallops during the spawning season, and (c) gonad index (GSI, gonad weight/soft tissue weight) from a nearby population during spawning season. Error bars are ± 1 SD. Full moons are indicated by open circles. Only gene copy numbers are reported in (b) due to the unknown cell concentration of the samples



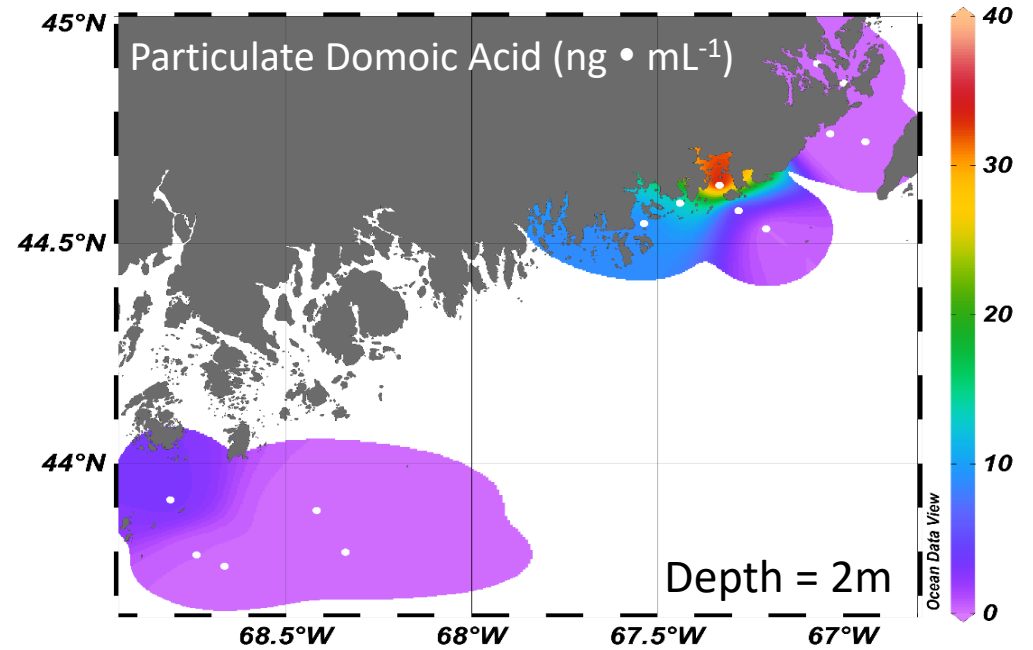
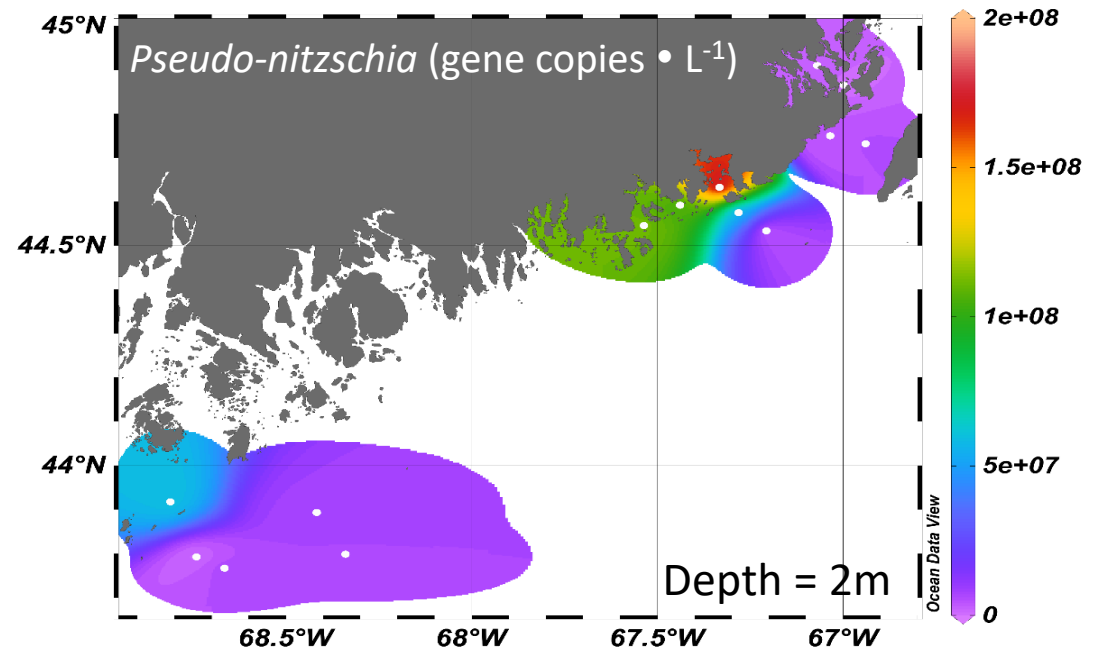
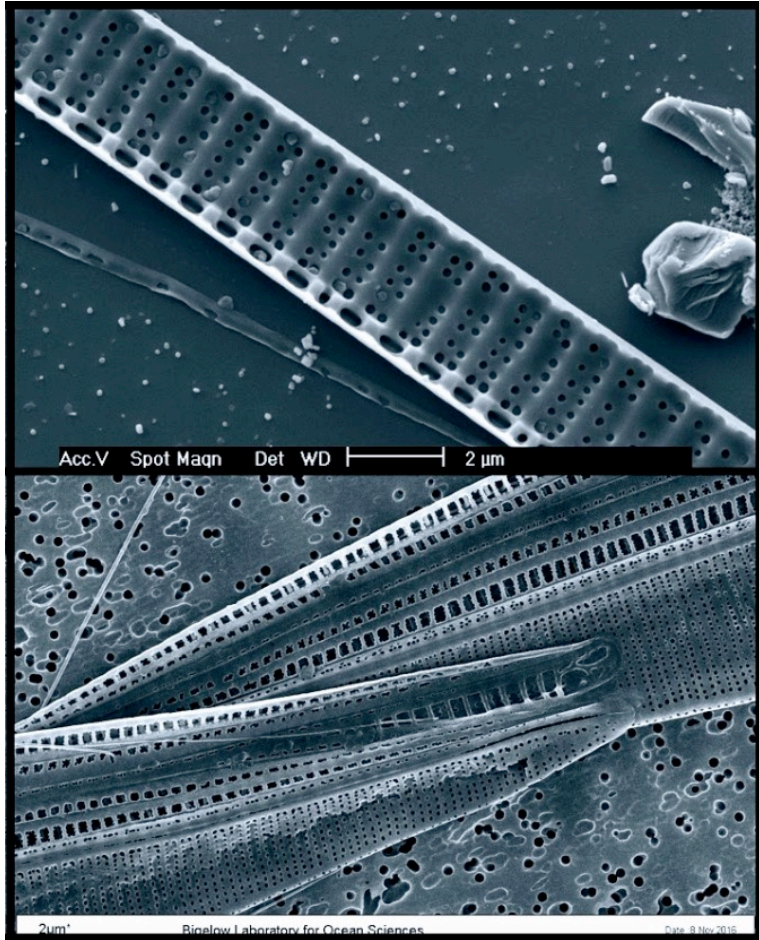
2. Portable, real-time eDNA

- Detects up to 3 gene targets in 9 samples
- Smart phone interface
- Uploads results to the cloud
- Can port existing assays directly
- 10 minute eDNA extraction kit (not shown here)



3. HABs and eDNA

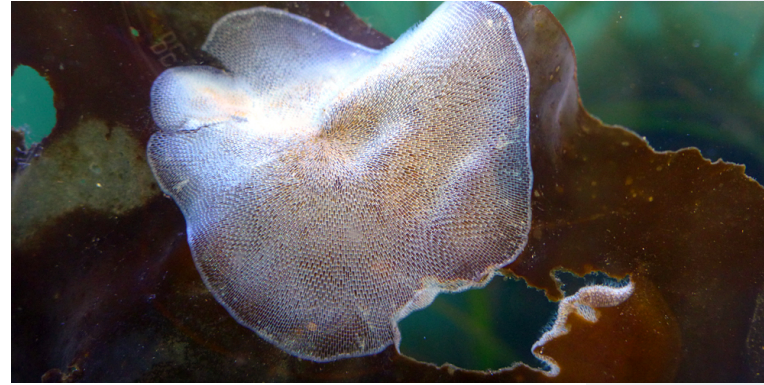
Scanning Electron Micrographs of *Pseudo-nitzschia* cells from the 2016 Bloom Event in the Gulf of Maine



Countway, Archer, Hubbard et al.

4. Comprehensive Site Selection Tool

- Eventually determine likelihood of lacey bryozoan or vibrio outbreak before choosing a site, *or* taking necessary steps to avoid closures or product loss



Thank You!

For More Information and Program Engagement:

<https://umaine.edu/eDNA/>

Supported by the National Science Foundation
under Grant No. 11A-1849227



And You!