

Welcome to this week's presentation and conversation hosted by the
Canadian Association for the Club of Rome,
a Club dedicated to intelligent debate and action on global issues.

Connectivity between freshwater and marine ecosystems.

Our speaker today is Dr. Robin Sleith, who is with the Maine environmental DNA project, a multi-institutional initiative that seeks to better understand Maine's coastal ecosystems. Using a combination of eDNA tools such as metabarcoding and quantitative PCR, we are testing ecological hypotheses around the boom-and-bust dynamics underlying harmful algal blooms in marine and freshwater ecosystems. Ultimately, we want to understand the factors that promote or impede formation of harmful algal blooms to forecast and manage these events better.

DESCRIPTION: This talk will explore the connectivity between freshwater and marine ecosystems, using the ongoing Maine eDNA project as a guide. Climate change and nutrient pollution in freshwater ecosystems have led to increased impacts on nearby marine ecosystems. Furthermore, anadromous species that rely on freshwater during their lifespan both impact and are impacted by freshwater systems. Emerging technologies like eDNA are able to increase the frequency and resolution of monitoring programs and help us understand these dynamic processes.

The presentation will be followed by a conversation, questions, and observations from the participants.

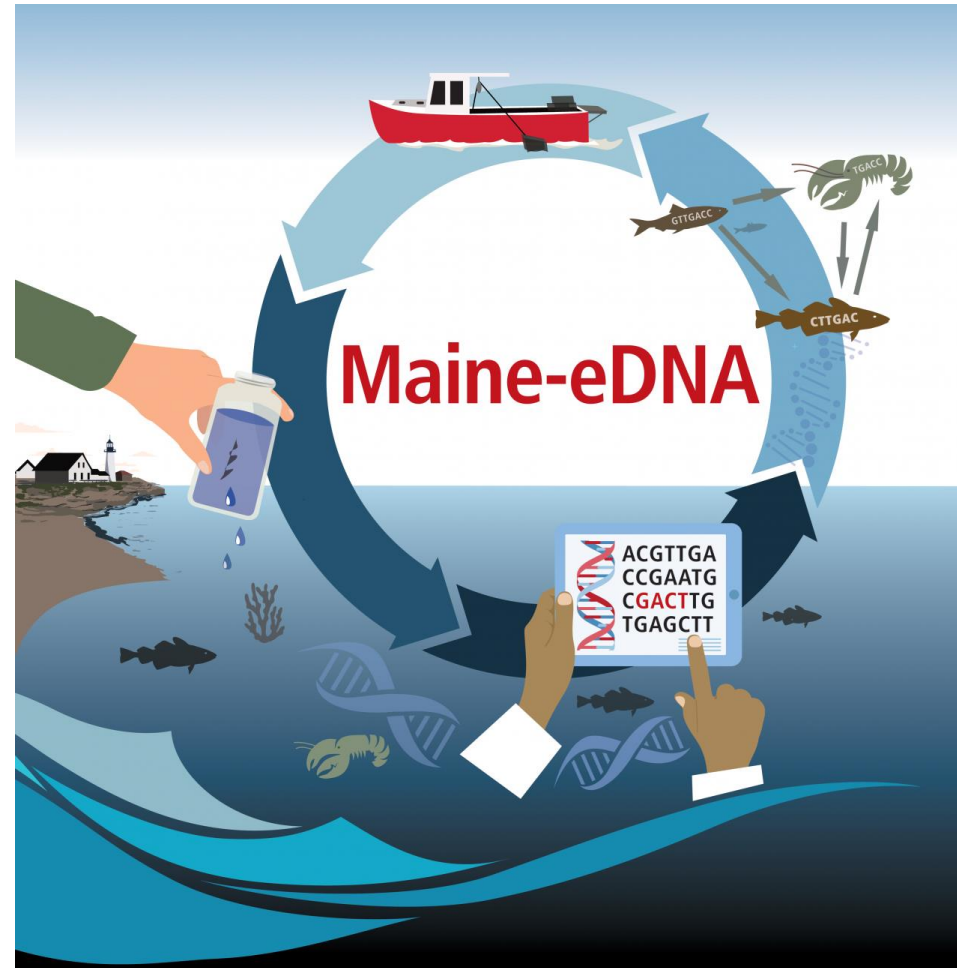
CACOR acknowledges that we all benefit from sharing the traditional territories of local Indigenous peoples (First Nations, Métis, and Inuit in Canada) and their descendants.



Website: canadiancor.com
Twitter: [@cacor1968](https://twitter.com/cacor1968)
YouTube: [Canadian Association for the Club of Rome](https://www.youtube.com/channel/UC...)
2023 May 24 Zoom #148

Maine-eDNA

- Molecule to Ecosystem: Environmental DNA as a Nexus of Coastal Ecosystem Sustainability for Maine
- A \$20 million NSF EPSCoR Research Infrastructure Improvement Track-1 Award



Genetic material obtained directly from environmental samples without any obvious signs of the biological source material (Thomsen and Willerslev, 2015)

Where does eDNA come from?

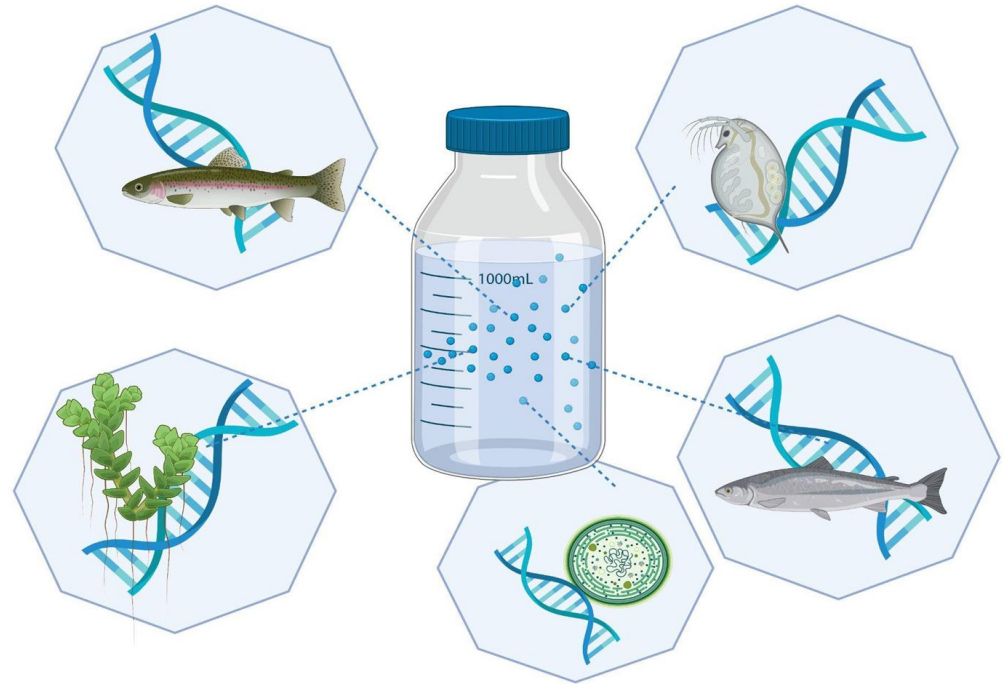
- Cellular decomposition
- Whole shed cells
- Whole microorganisms

Where is eDNA found?

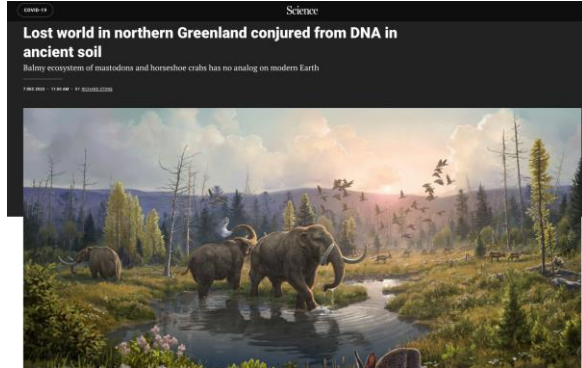
- Water
- Soils
- Air

How is eDNA used?

- Community characterization (metabarcoding)
- Targeted detection & quantification (qPCR)



eDNA in the news



The New York Times

Your DNA Can Now Be Pulled From Thin Air. Privacy Experts Are Worried.

Environmental DNA research has aided conservation, but scientists say its ability to glean information about human populations and individuals poses dangers.



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SCIENCE

LISTEN & FOLLOW

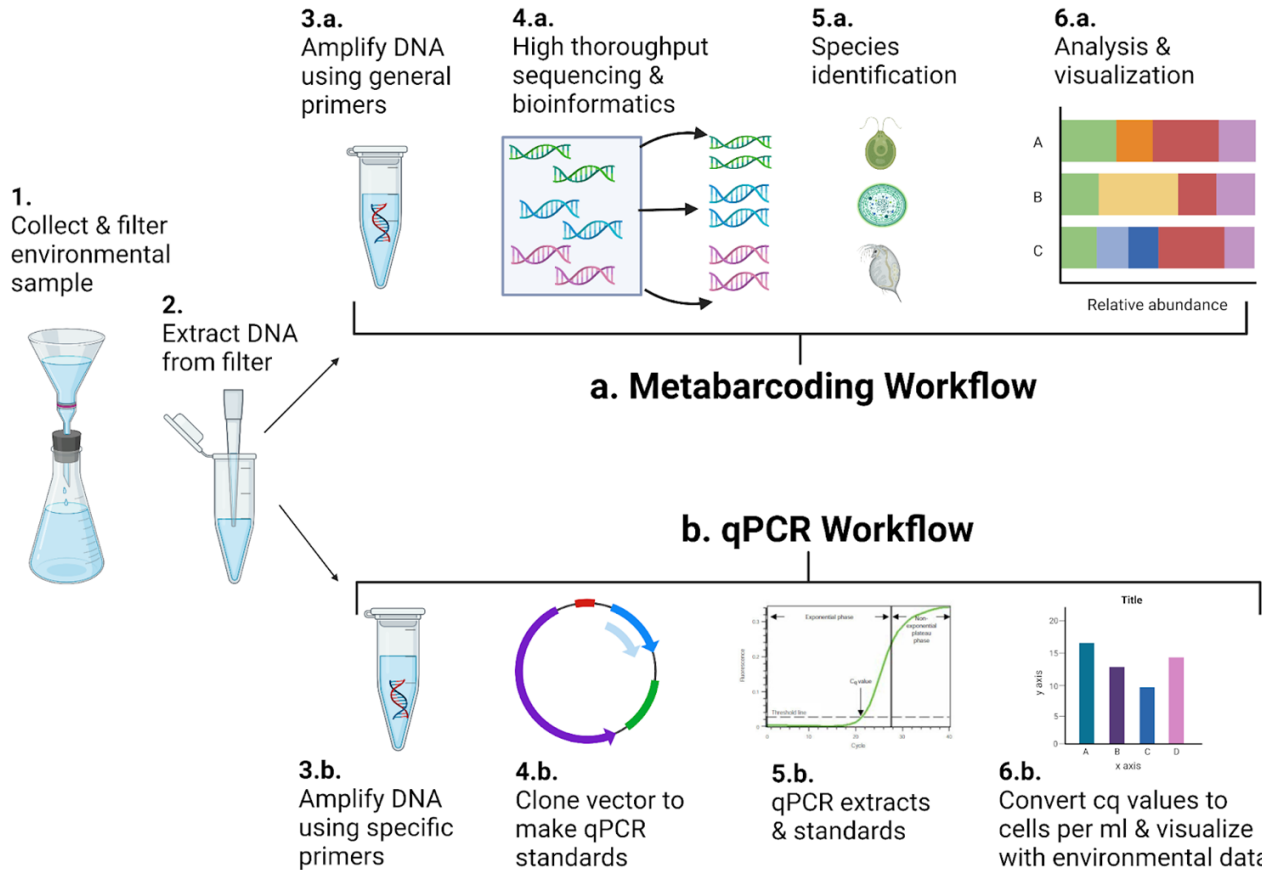
Scientists vacuum zoo animals' DNA out of the air

January 9, 2022 · 7:00 AM ET

Geoff Brumfiel



Researchers were able to detect DNA from elephants at the Copenhagen Zoo simply by sampling the air nearby.
by Marie Ojgaard /Ritzau Scanpix/AFP via Getty Images



RESEARCH

Goal 1: Sustainable Fisheries

Goal 1.1
Ecosystem Restoration

Goal 1.2
Larval Black Box

Goal 2: Harmful Species

Goal 2.1
Harmful Blooms

Goal 2.2
Species on the Move

Goal 3: Macrosystem eDNA Integration

Goal 3.1
Big Data Integration: Dynamics & Stability

Goal 3.2
Microbial Communities as Biosensors of Change

Goal 3.3
Comparative Study of eDNA Team Science

PROJECT ROADMAP



PROJECT ELEMENTS

Goal 4.1
Education and Workforce Development

Goal 4.2
Seed Funding and Emerging Areas

Goal 4.3 Diversity

Goal 4.4
Partnerships and Collaboration

Goal 4.5
Communication and Dissemination

Goal 4.6 Sustainability

Goal 4.7
Management, Evaluation, and Assessment



Harmful Species Vignette 1: Lakes, Cyanobacteria, and Oysters

What is a HAB?

- “Out of control” growth of algae
- Impacts many sectors
- Commercial and recreational fisheries
- Aquaculture
- Tourism
- Wildlife
- Human health

HARMFUL ALGAL BLOOMS

Harmful algal blooms, or HABS, occur when colonies of microscopic algae grow out of control. These blooms are a growing problem in every U.S. coastal and Great Lakes state. While we can't prevent these blooms, we can be better prepared. NOAA leads many research efforts to help coastal communities counter the environmental and health effects associated with these "red tide" events.

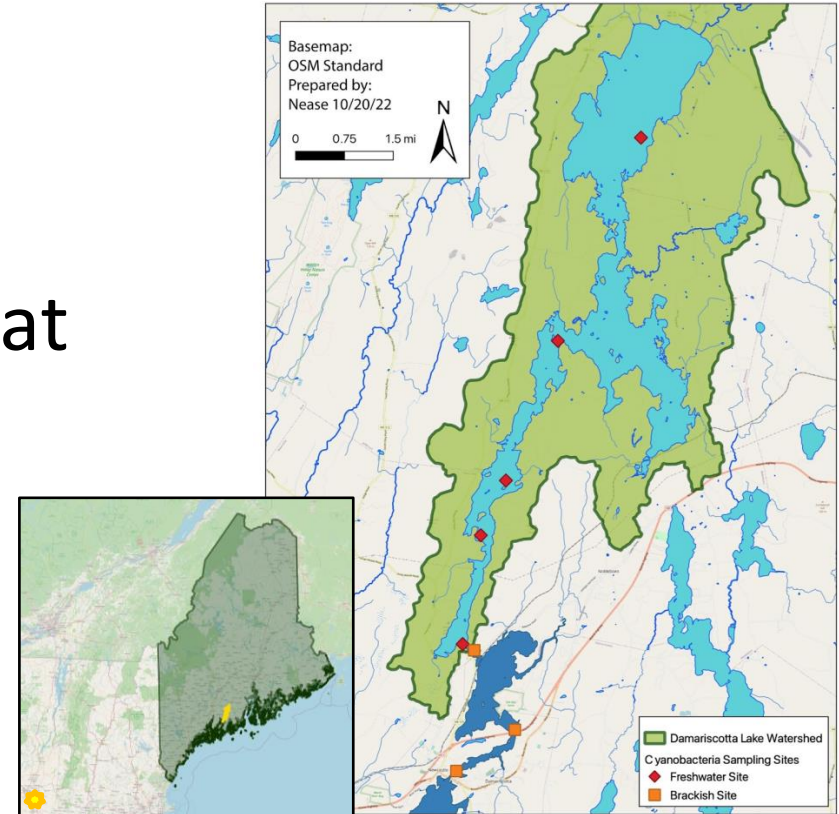
The infographic features four circular icons illustrating the impacts of HABs: a water tap with a red drop (human health), a dead bird (wildlife), a red shellfish (aquaculture/fisheries), and a fish with a yellow and green pattern (tourism/recreation).

Sometimes, microscopic algal species in waterways around the nation grow out of control. Some of these algal blooms can contaminate water and shellfish, kill animals, and make humans sick.

NOAA

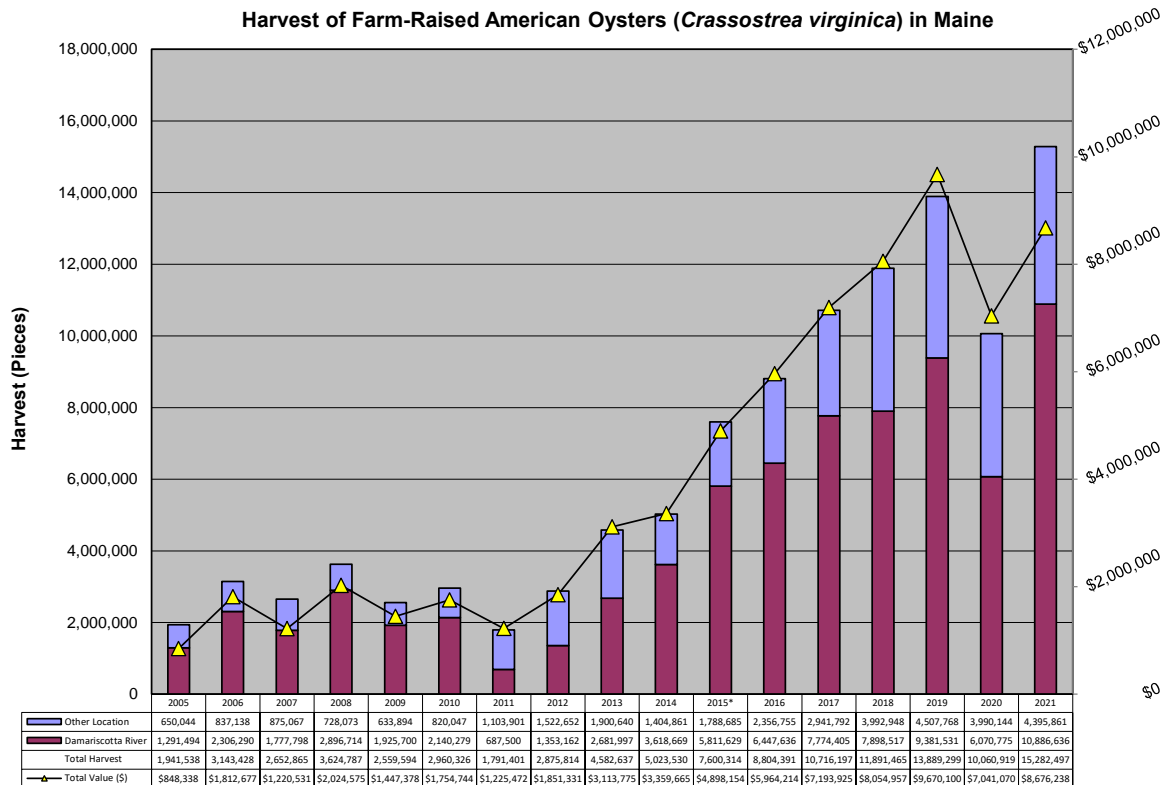
Damariscotta Lake Ecosystem

- 4,600 acres
- 22 miles of shoreline
- State park & public boat launch
- 3 distinct basins
- Large alewife run (>1 million fish!)



Damariscotta River Estuary

Harvest of Farm-Raised American Oysters (*Crassostrea virginica*) in Maine

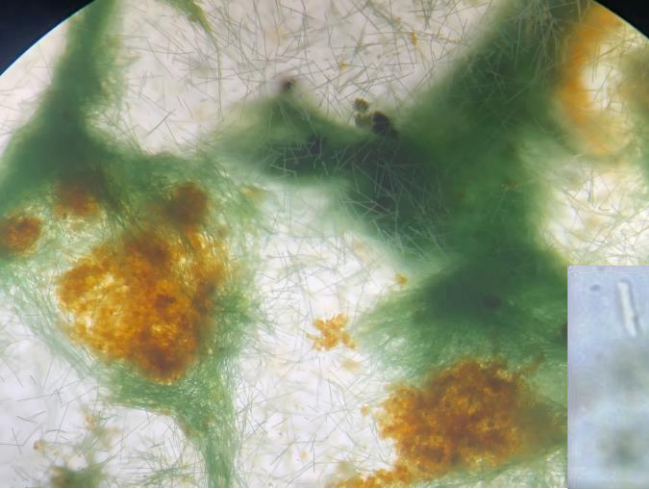


* DMR began collecting LPA harvest data in 2015.

* 2021 data are preliminary and will be finalized when 2022 data are posted.

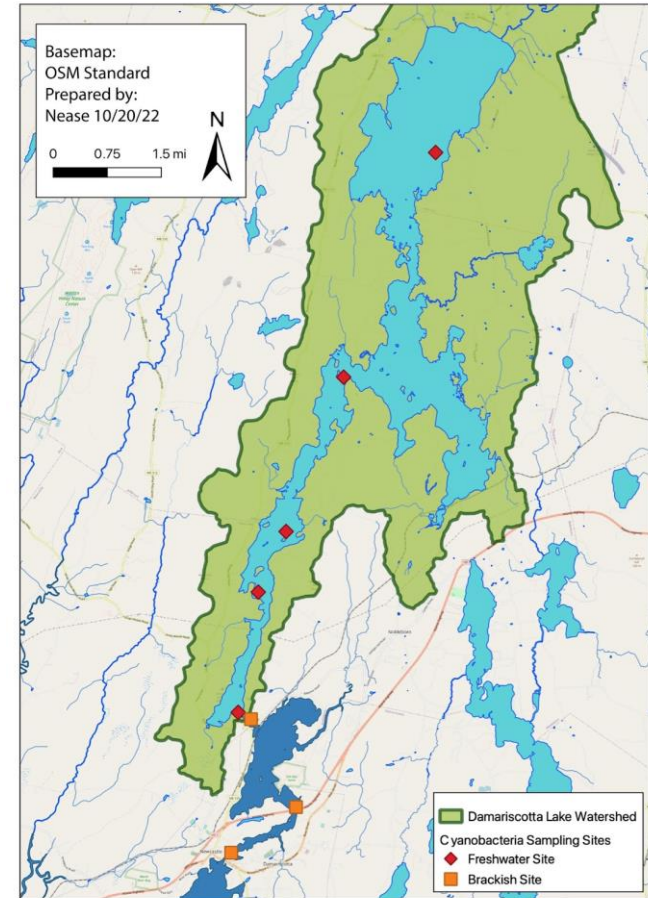


2020 *Planktothrix* Bloom

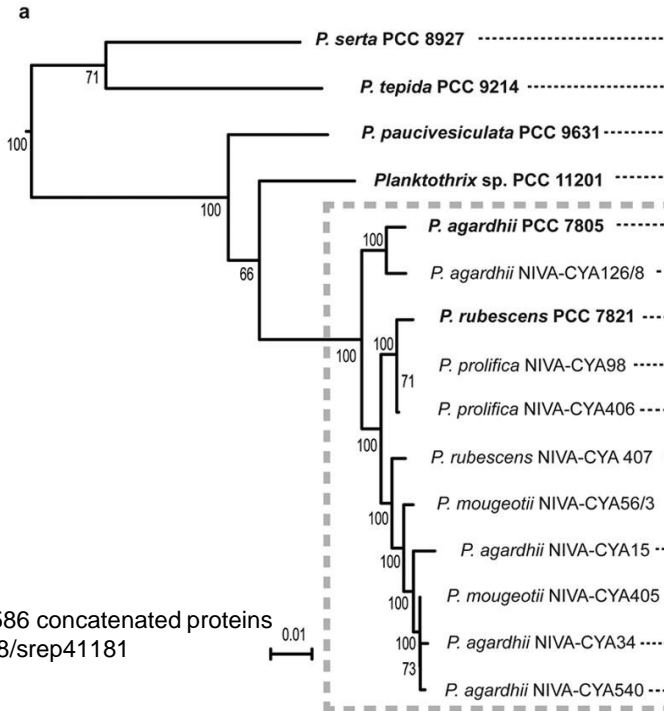


Research Questions

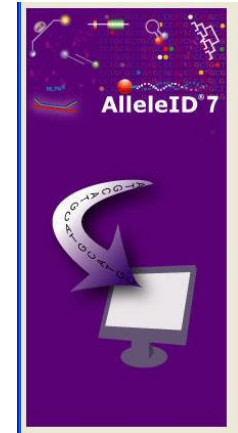
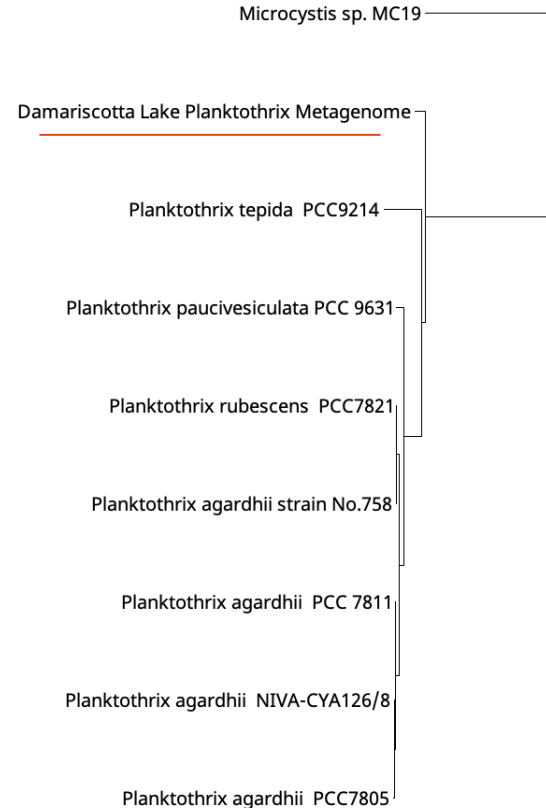
- What is the distribution and duration of the bloom
- Toxicity?
- Are cells/toxins moving downstream?
- What factors (biotic and abiotic) are associated with bloom formation and cessation



Planktothrix primer development

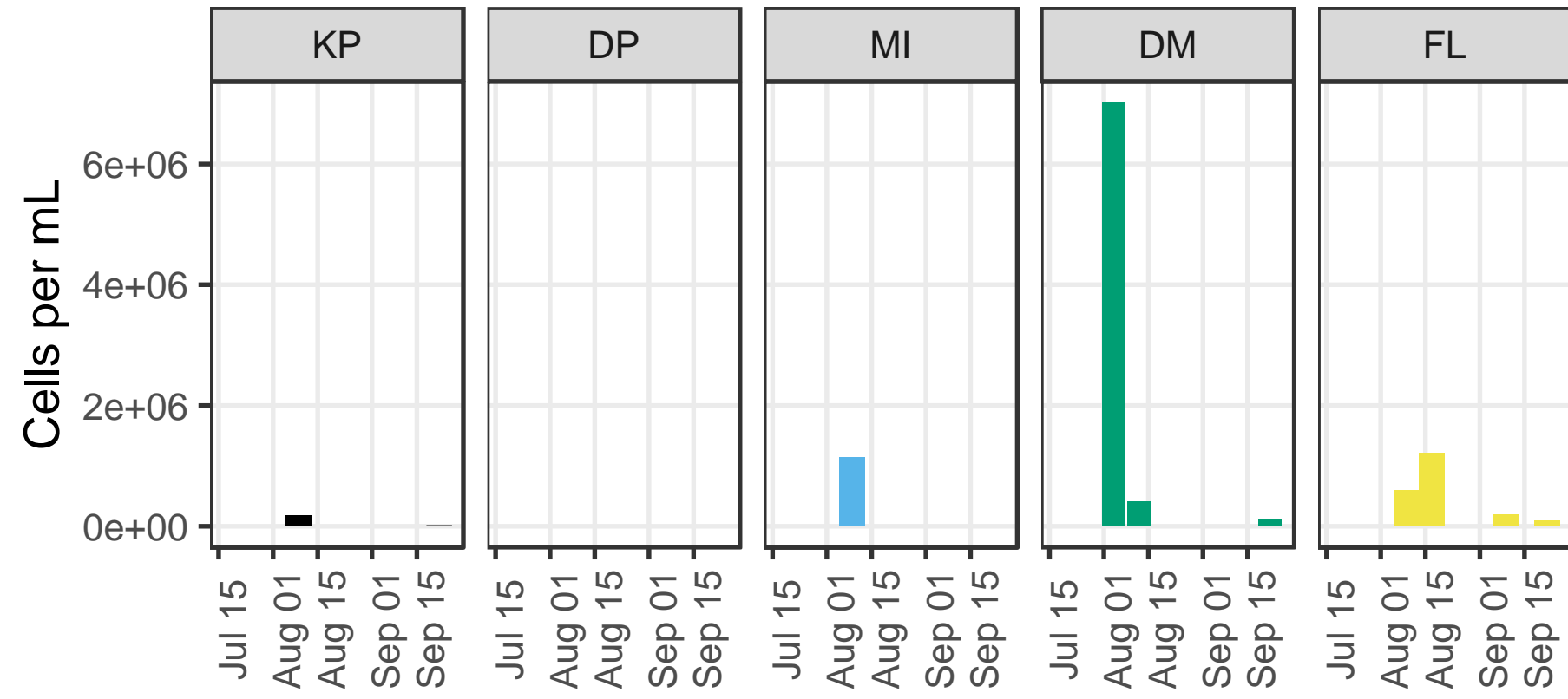


Panrace et al 2017 586 concatenated proteins
<https://doi.org/10.1038/srep41181>



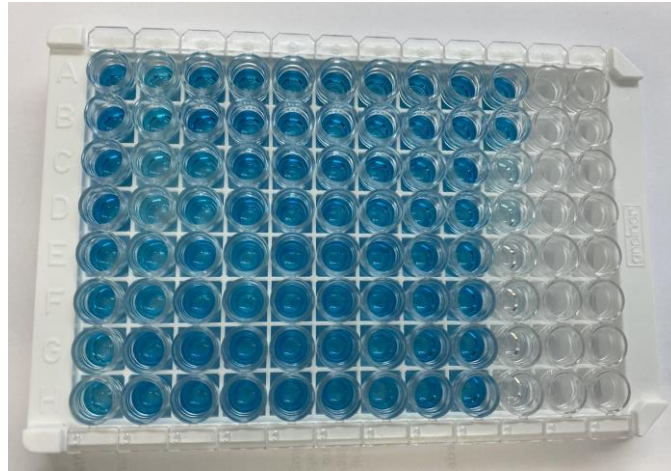
16S, ITS, 23S Alignment

2021 *Planktothrix* qPCR results



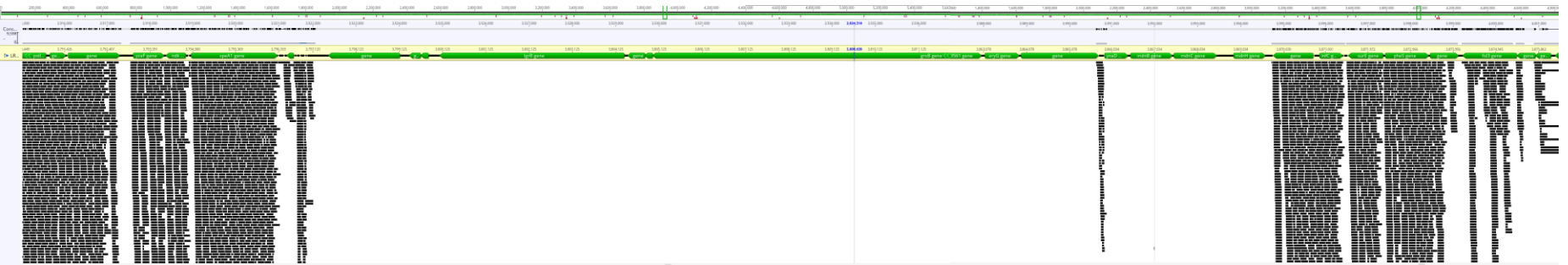
Planktothrix ELISA results

- Abraxis Microcystins/Nodularins (ADDA) kit
- No results above lowest threshold (0.3 ug/L)



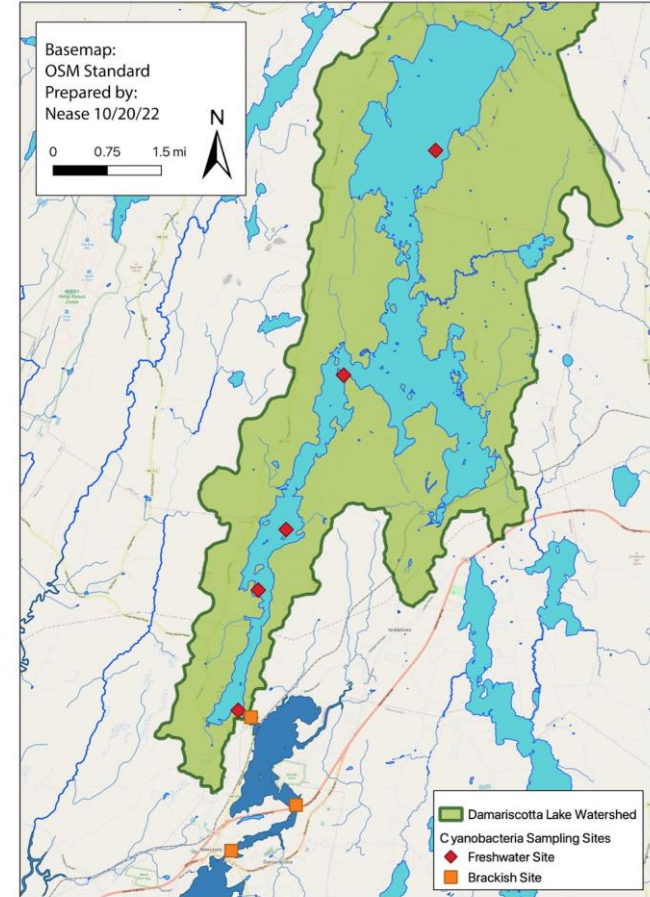
Planktothrix microcystin biosynthesis

- Damariscotta Lake *Planktothrix* lacks microcystin biosynthesis pathway

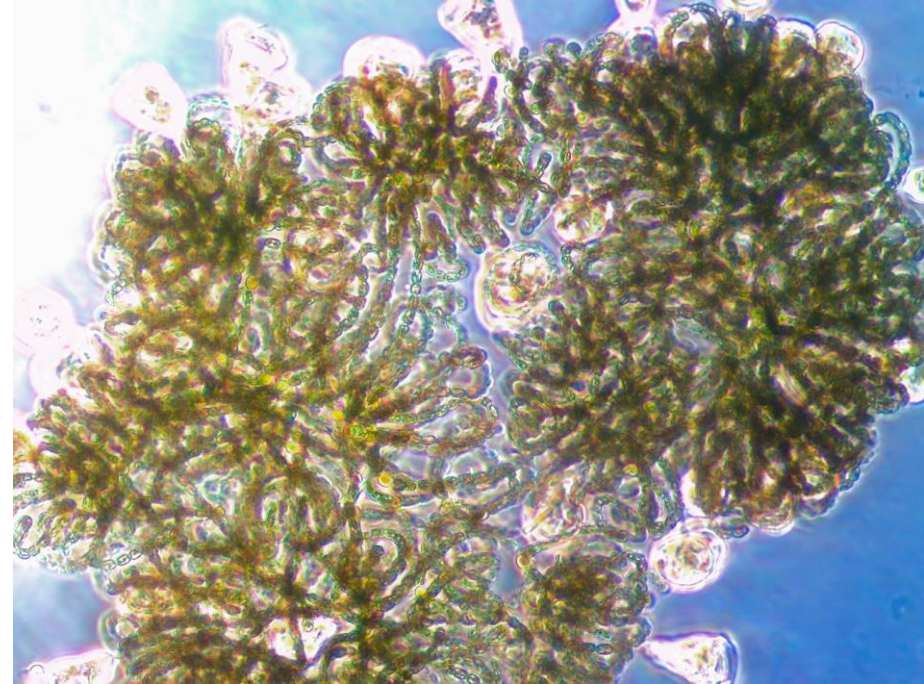
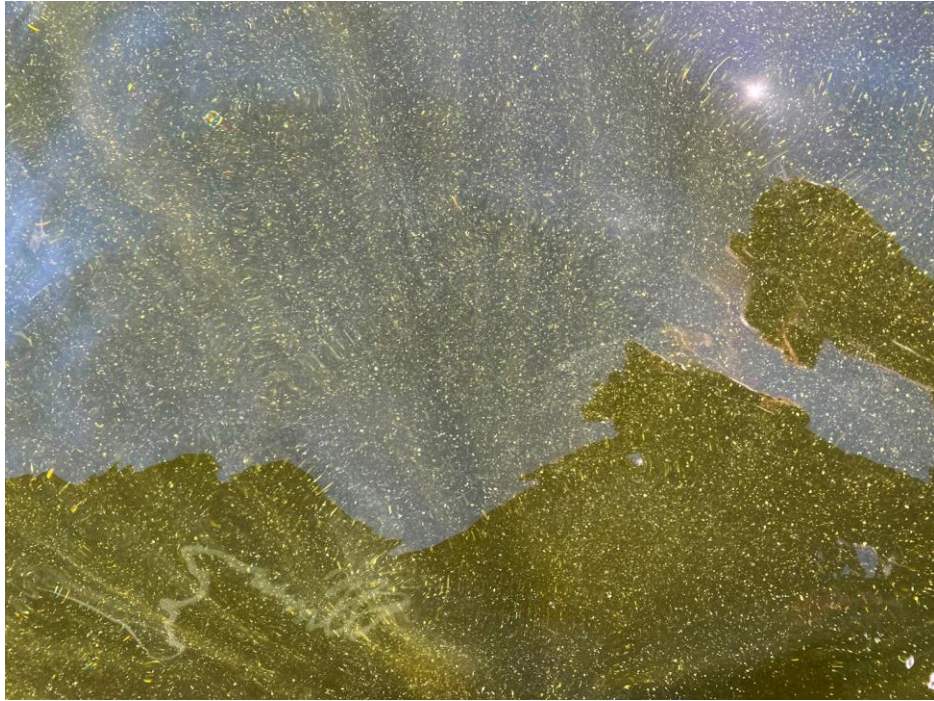


Plans for 2022

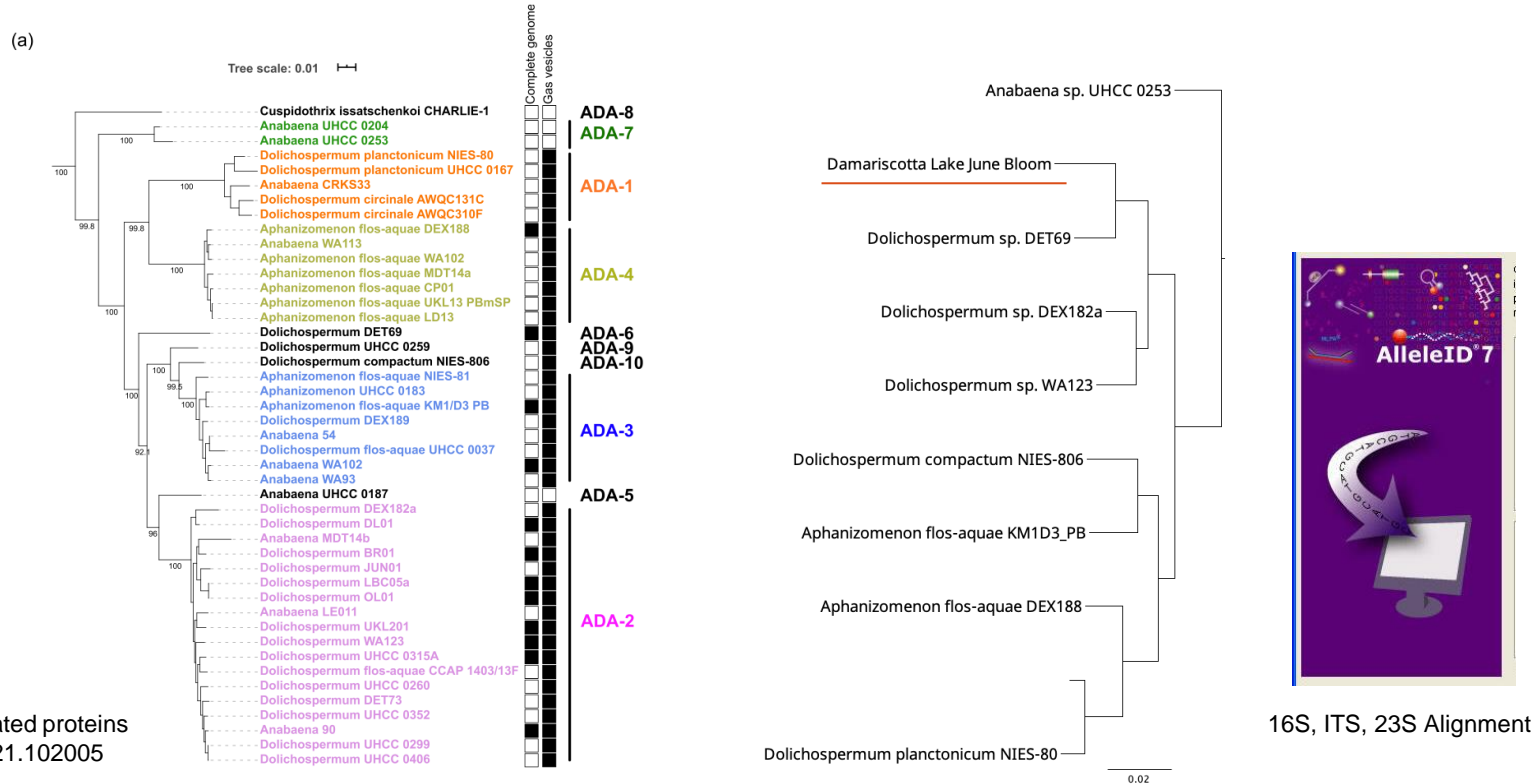
- Expand sampling further downstream
- Expand sampling window (June-October)



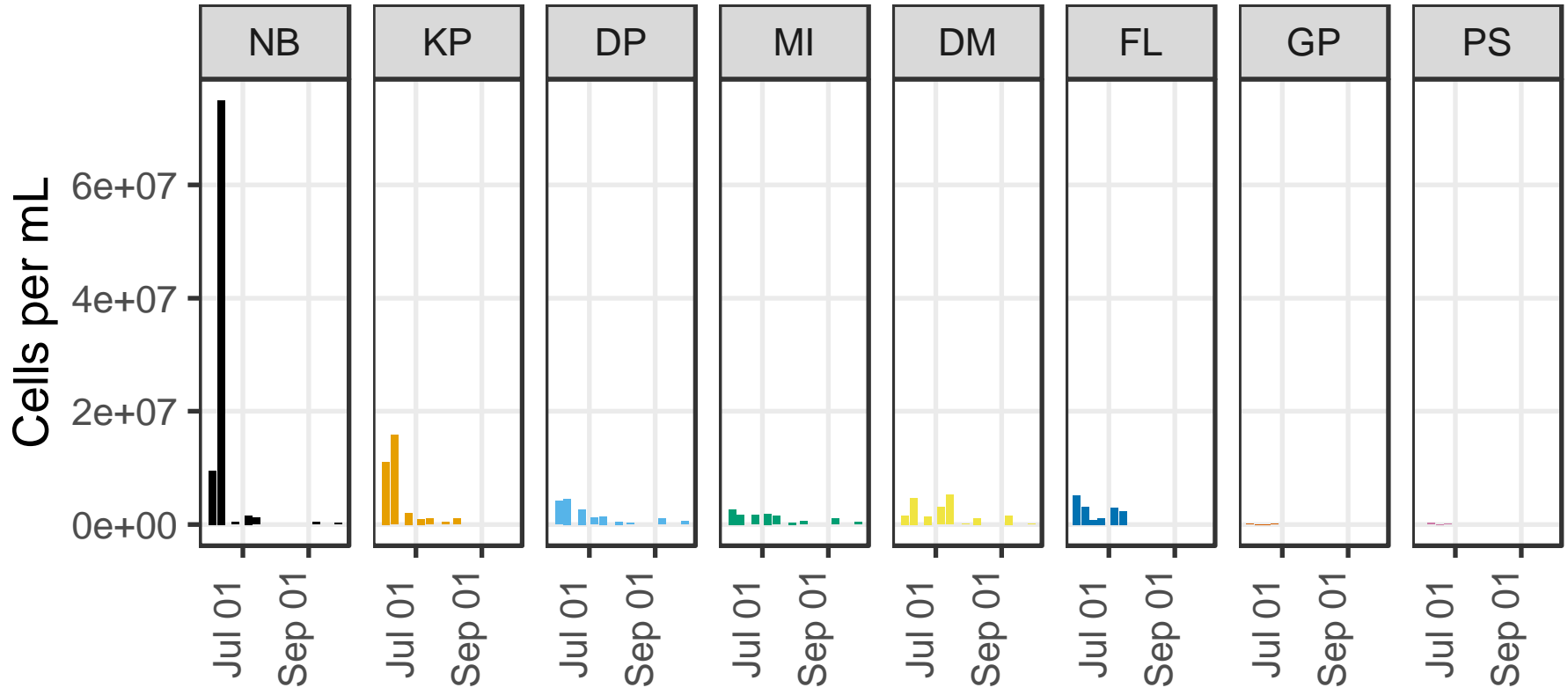
June *Dolichospermum*



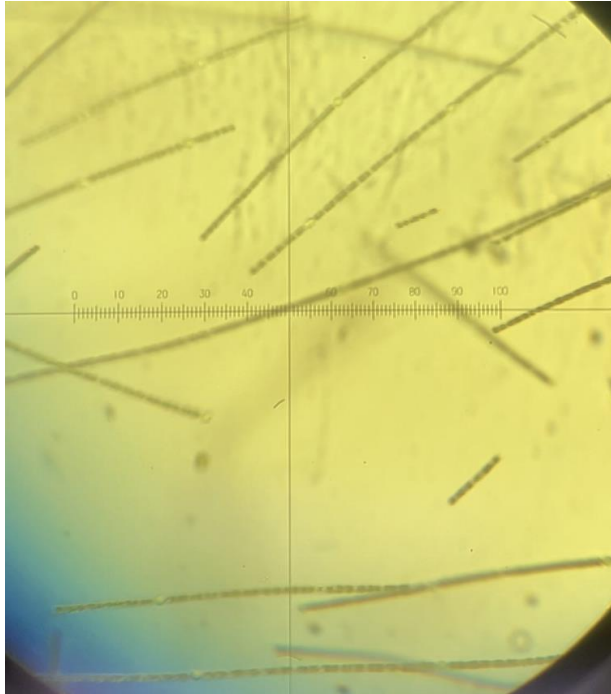
Dolichospermum primer development



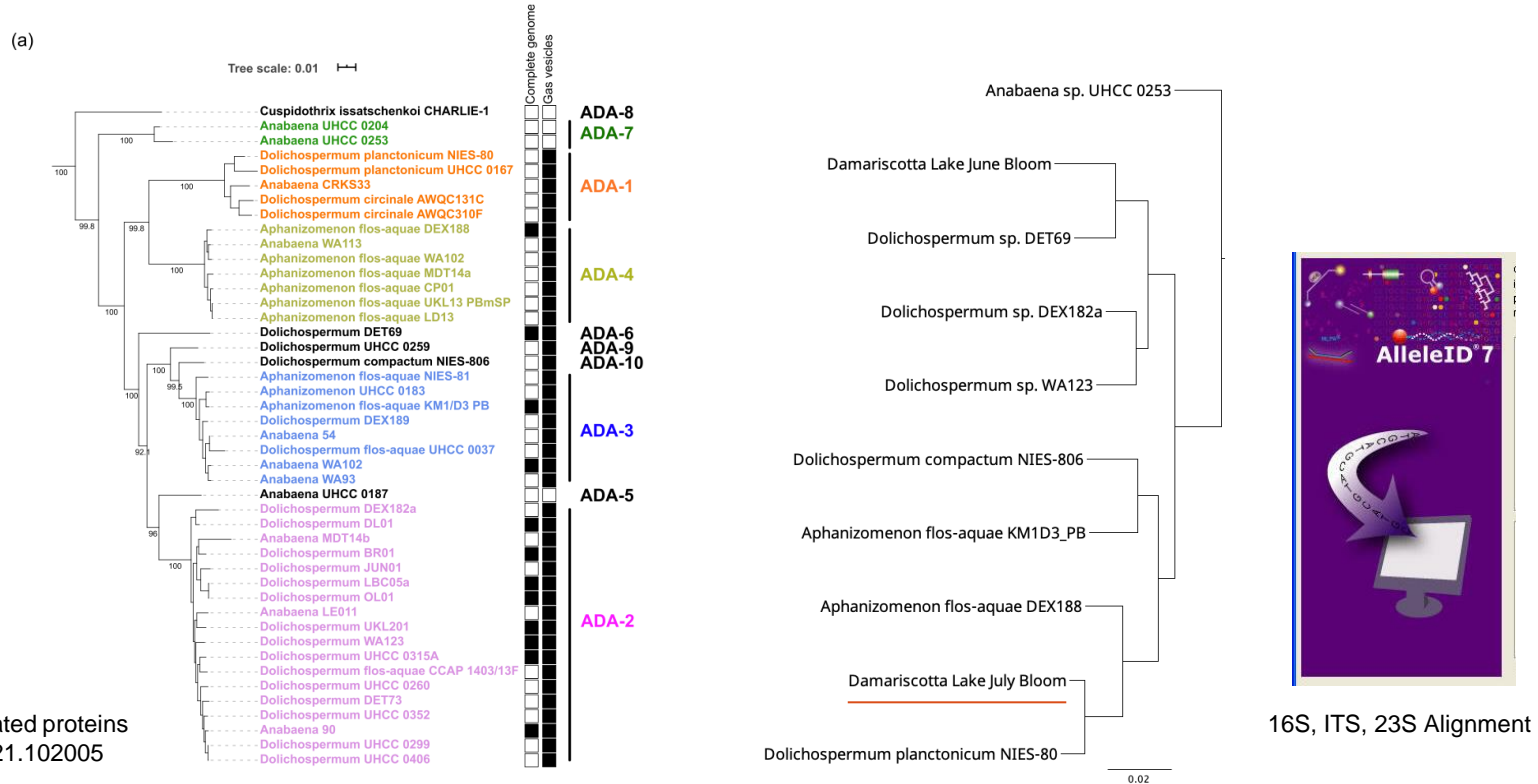
June *Dolichospermum* ddPCR counts



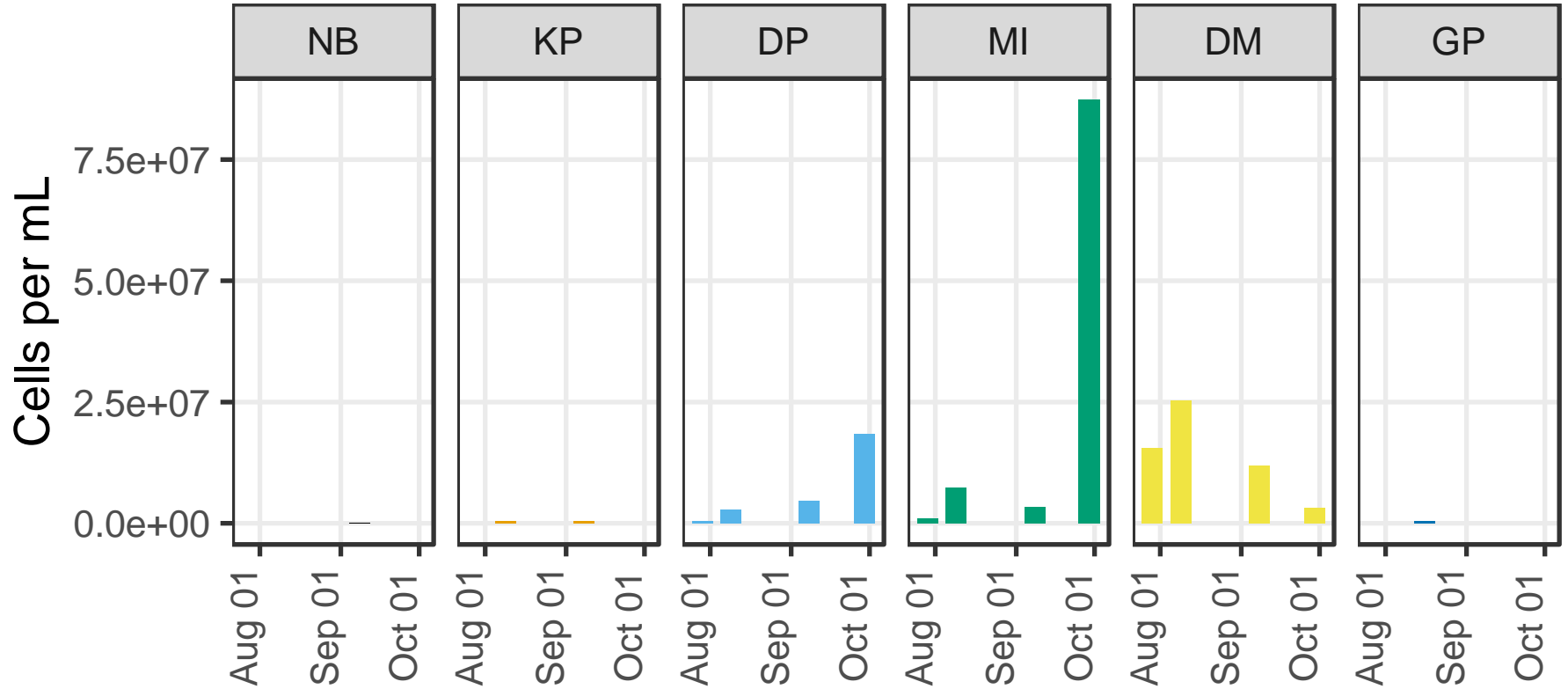
The Sequel: July *Dolichospermum*!



July *Dolichospermum* primer development



July *Dolichospermum* ddPCR counts



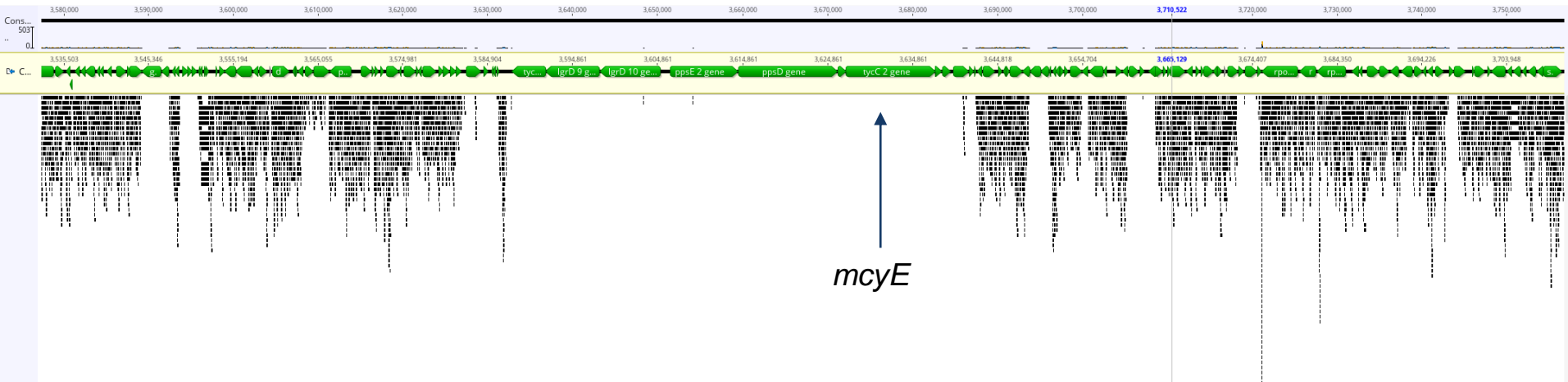
Dolichospermum metagenomes

- 66% Nostocales
- Antismash analysis
 - Anabaenopeptin
 - Cylindrospermopsin
 - Cyanobactin



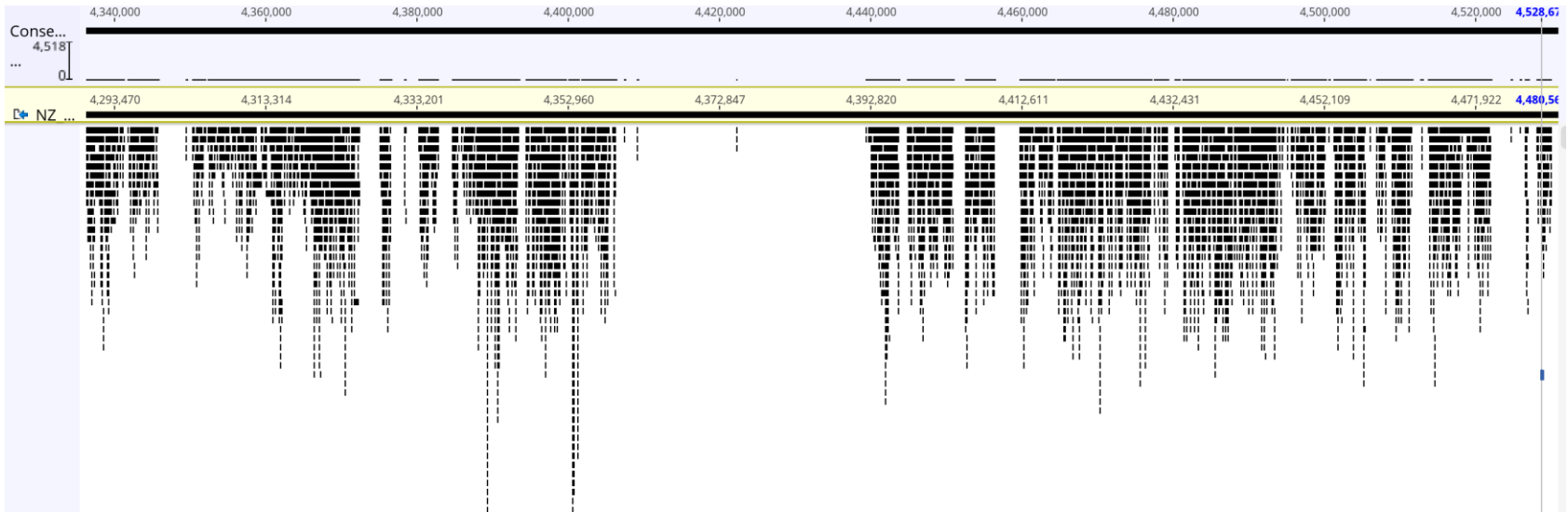
Dolichospermum microcystin biosynthesis

- Damariscotta Lake *Dolichospermum* lack microcystin biosynthesis pathway



Dolichospermum anatoxin biosynthesis

- Damariscotta Lake *Dolichospermum* lack anatoxin biosynthesis pathway



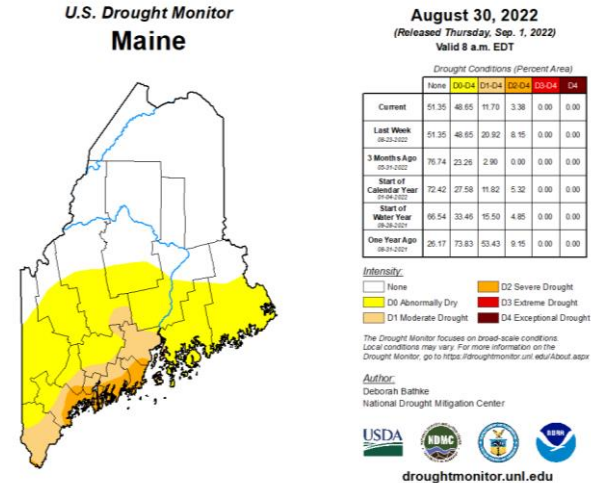
Don't forget about *Planktothrix*!

- Downgrading to nuisance bloom (for now)
- Restricted to upper reaches of South Basin in July



Conclusions and Next Steps

- Did a dry 2022 summer favor *Dolichospermum*?
- No evidence of survival in brackish water
- Unable to characterize microcystin transport (no microcystin detected)
- Incorporate nutrient and environmental data
- *mcyE* containing *Dolichospermum* genotype found 20 miles north, is it a threat?
- 2023 sampling oyster tissues just downstream for microcystin and anabaenopeptin



Precedent

- Freshwater *Microcystis* bloom
- Downstream transport
- Bioaccumulation of microcystin in shellfish
 - Up to 107x higher than adjacent seawater
- Acute liver failure for otters consuming shellfish

Sea otter deaths linked to toxin from freshwater bacteria

September 10, 2010

By Tim Stephens

A potent toxin produced by bright-green blooms of freshwater bacteria has been flowing into the ocean and poisoning sea otters, according to a team of investigators led by scientists at the California Department of Fish and Game (DFG) and the University of California, Santa Cruz.

In a paper published September 10 in *PLoS ONE*, the researchers reported that the deaths of at least 21 California sea otters (a federally listed threatened species) were linked to the toxin, called microcystin. First author Melissa Miller, a senior wildlife veterinarian at the DFG Marine Wildlife Veterinary Care and Research Center in Santa Cruz, will present the findings at the California and the World Ocean Conference in San Francisco on Friday, September 10.

The toxin is produced by a type of cyanobacteria (also known as "blue-green algae") called *Microcystis*. Warm, nutrient-rich water conditions favor large blooms of *Microcystis*, which can form thick green mats on the surface of the water.

"This study is significant because it is the first to establish a connection between freshwater contamination by microcystin and marine mammal mortality," said Miller, who is also affiliated with UCSC's Institute of Marine Sciences. "This land-to-sea link has important implications for marine life and human health."

Coauthor Raphael Kudela, professor of ocean sciences at UC Santa Cruz, said the team found high concentrations of microcystin in lakes bordering Monterey Bay and in rivers that flow into the bay.

"The toxin itself is extremely stable. In laboratory studies, we found that microcystin didn't degrade much even after three weeks in ocean water," Kudela said.

Microcystin poisoning can cause acute liver failure or damage other tissues and can be fatal. In 2007, Miller began seeing dead and dying sea otters recovered along the shore of Monterey Bay with evidence of acute liver failure. After tissue from the otters tested positive for microcystin, Miller teamed up with Kudela and DFG chemists to search for environmental

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The toxic cyanobacteria *Microcystis* gives this water sample from Pinto Lake its bright green color. (Photo by R. Ketley)

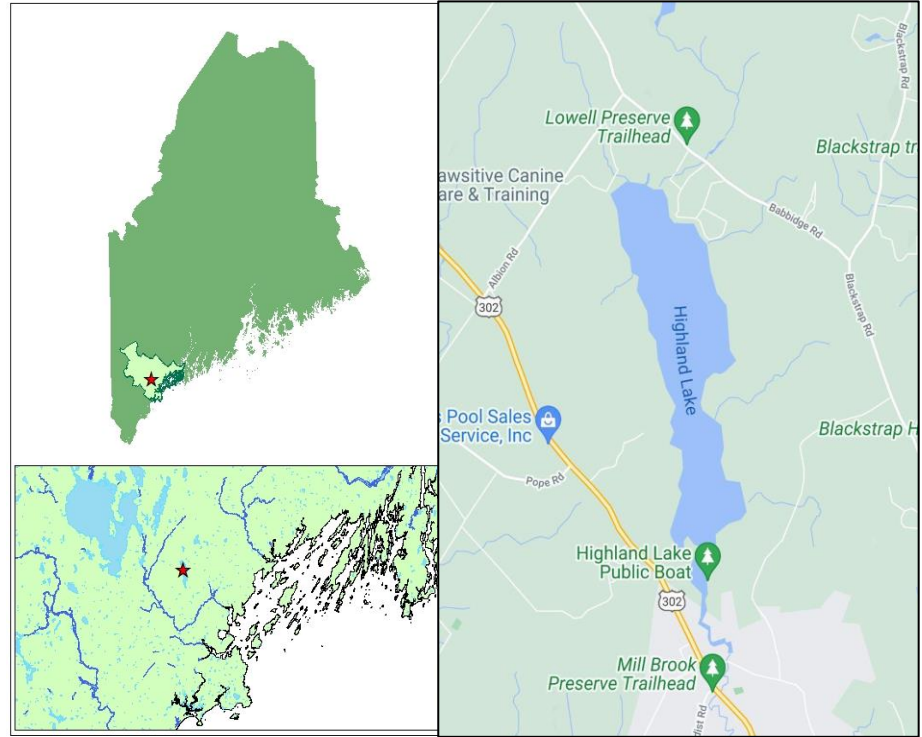


California sea otters have been poisoned by the microcystin toxin. This sea otter pup is healthy. (Photo by R. Wilder, Monterey Bay Aquarium)



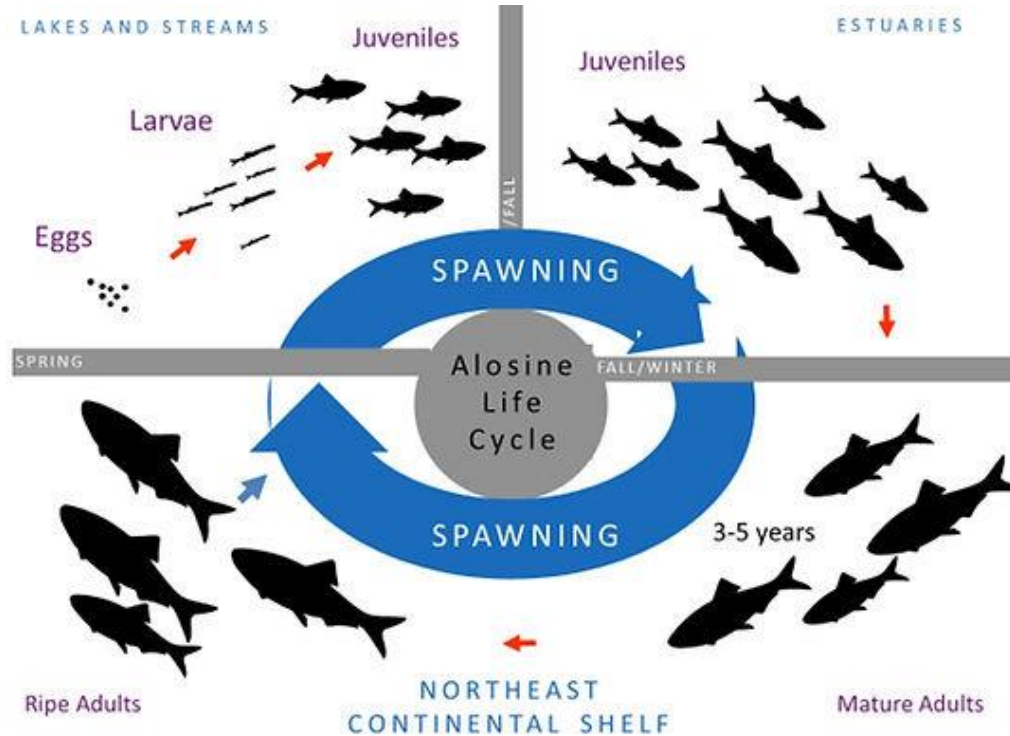
Harmful Species Vignette 2: Lakes, Algae, and Alewives

- Highly developed area
- Previously listed (1998-2010) as impaired by Maine-DEP due to deteriorating trophic state
- Active Lake Association water quality team
- Experienced an unusual nuisance bloom, coinciding with the first 4 years of high numbers of spawning anadromous alewife



Highland (Duck) Pond

Alewife life cycle



Cause(s) of algae bloom?

Excess nutrients?

Unusual phytoplankton species?

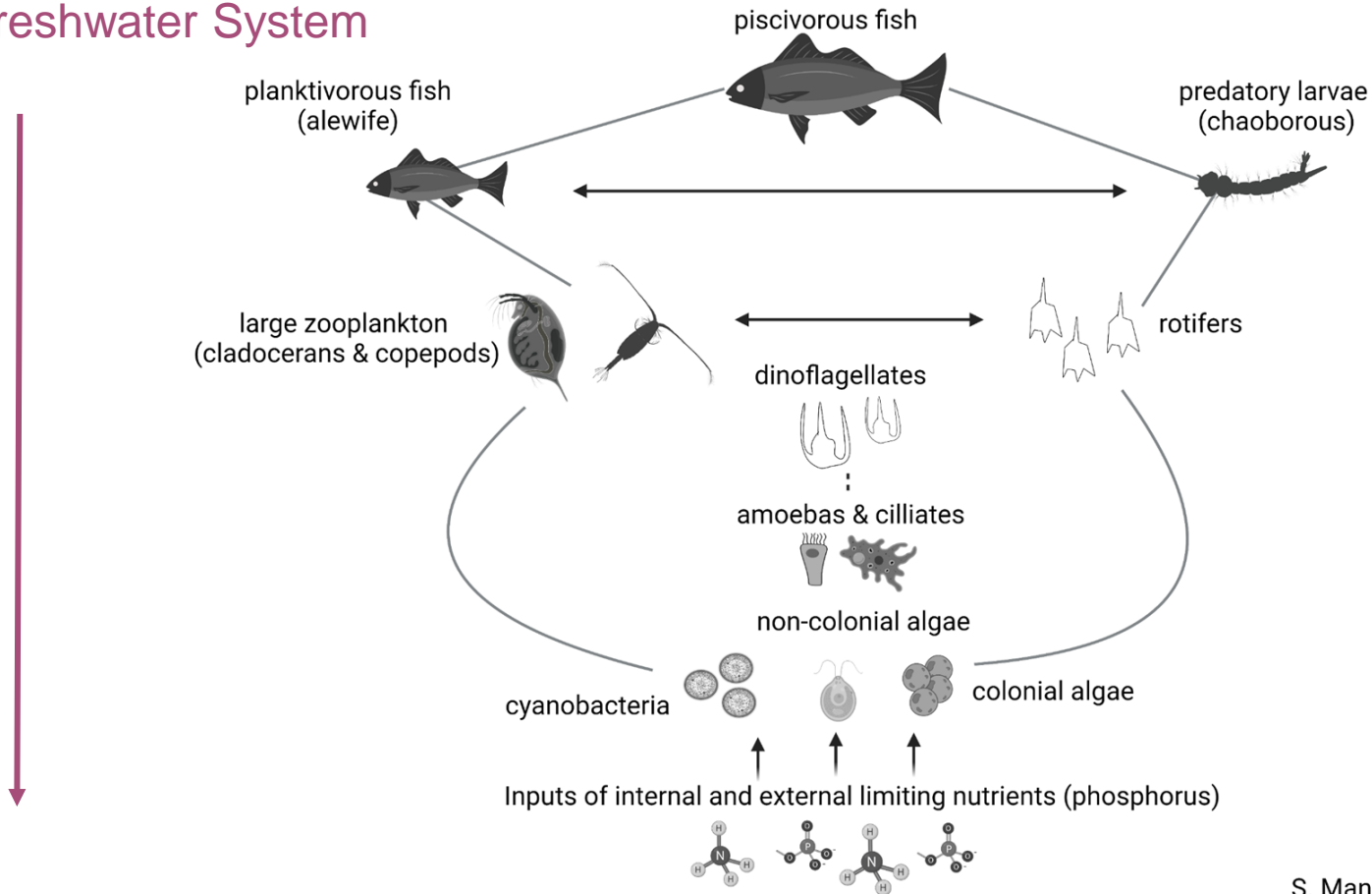
Trophic cascade triggered by consumption of herbivorous zooplankton by alewife?

Unlikely: TP levels lower than those associated with nuisance blooms

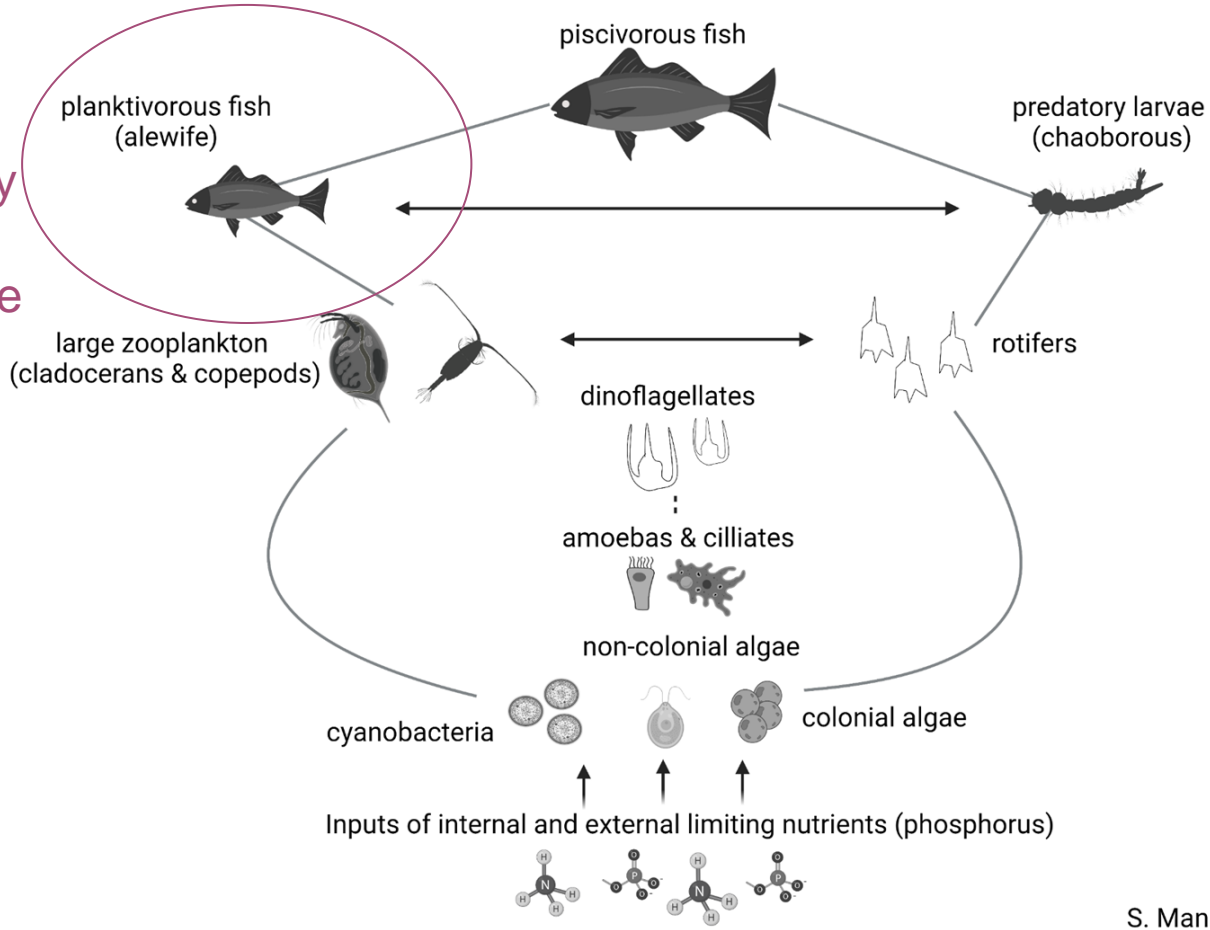
Preliminary observations suggested that the bloom was caused by a picocyanobacteria - Today's talk!

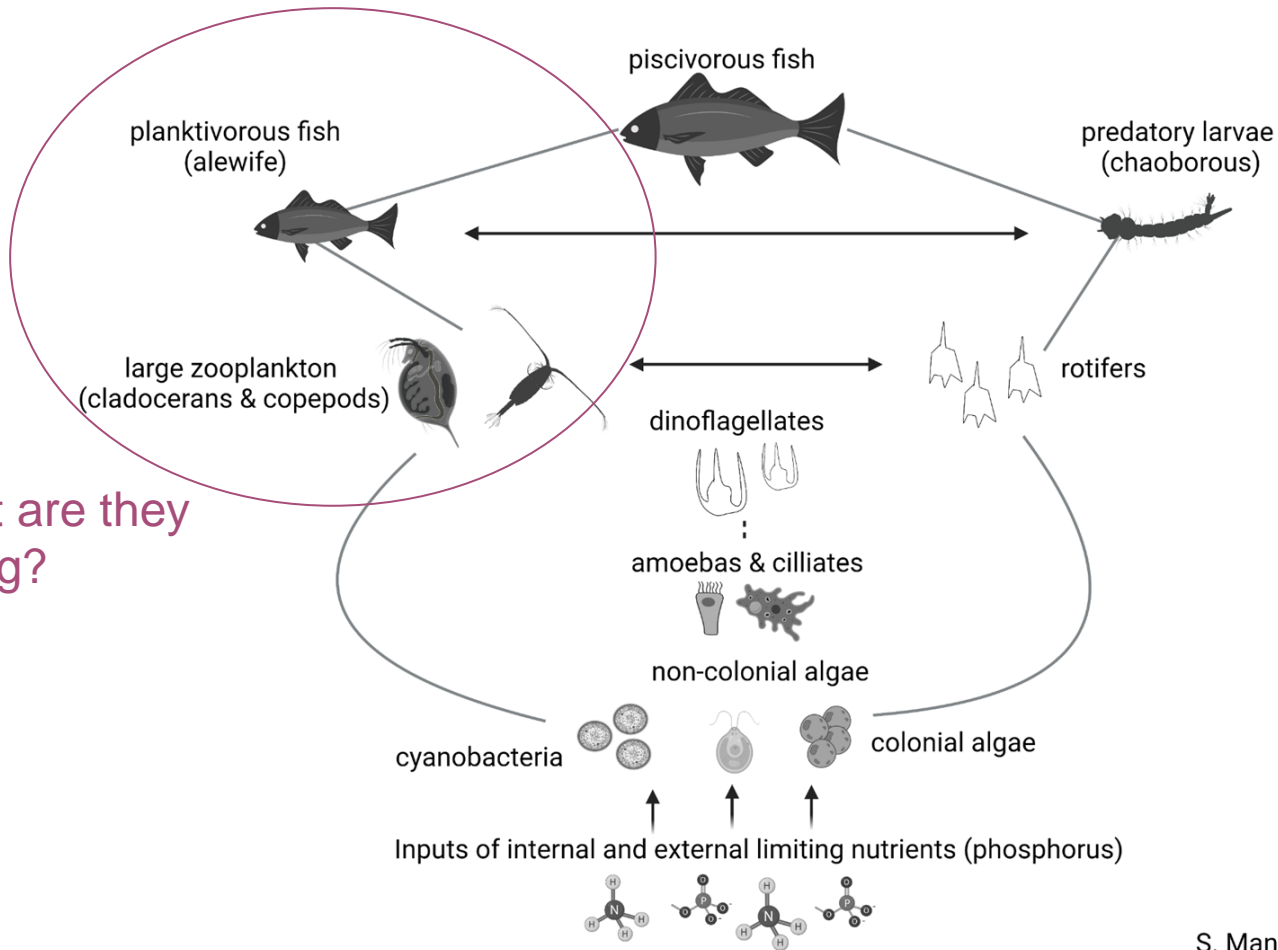
Coming soon!

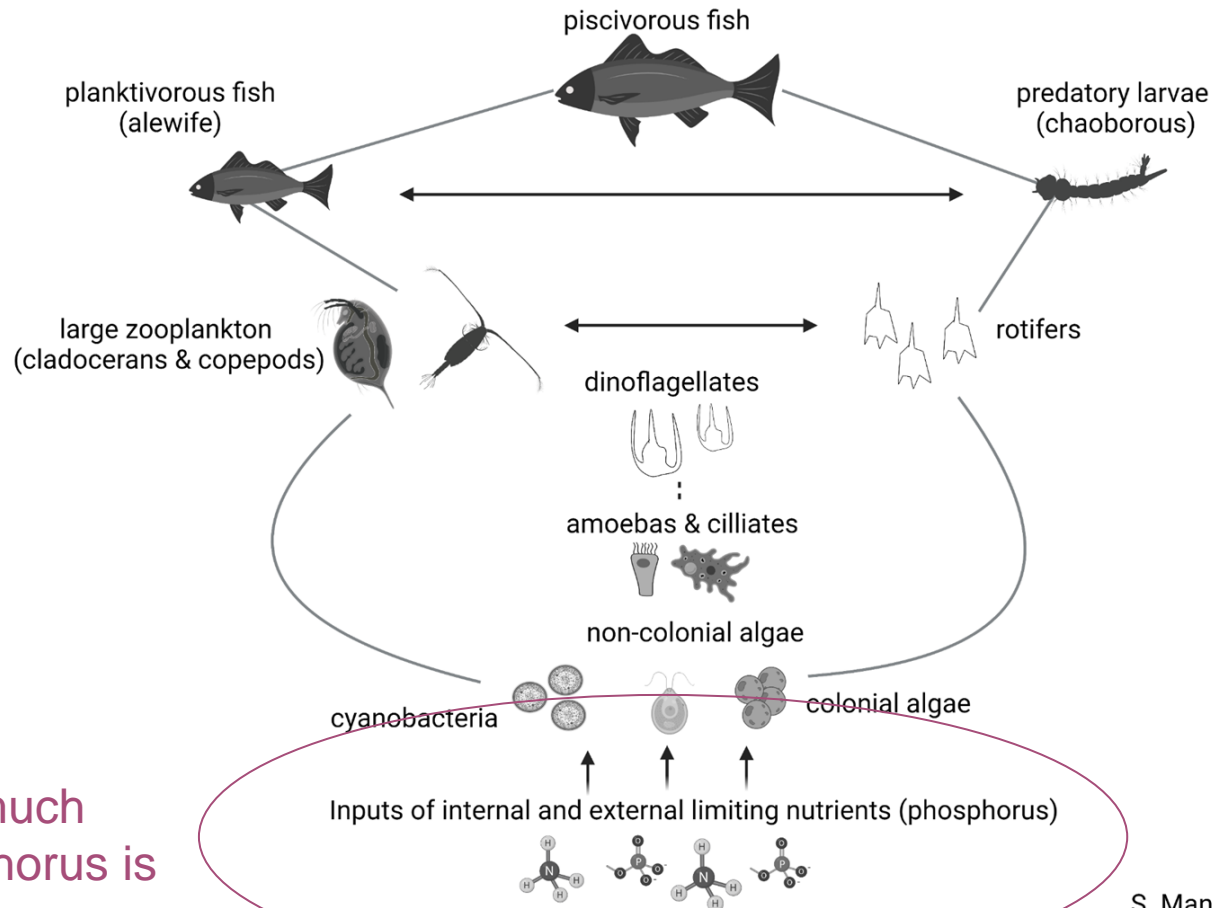
Freshwater System



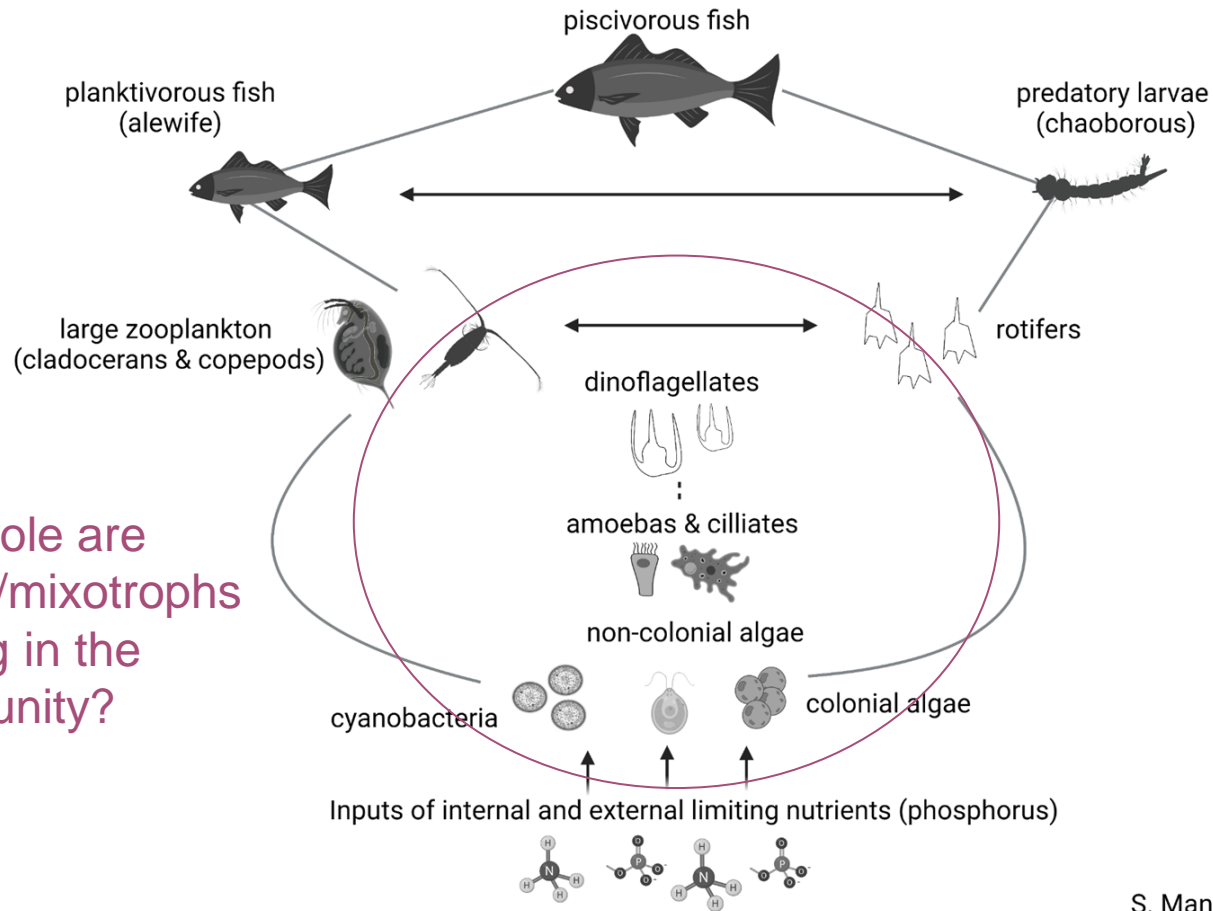
How many juvenile alewife are there?







How much phosphorus is there?



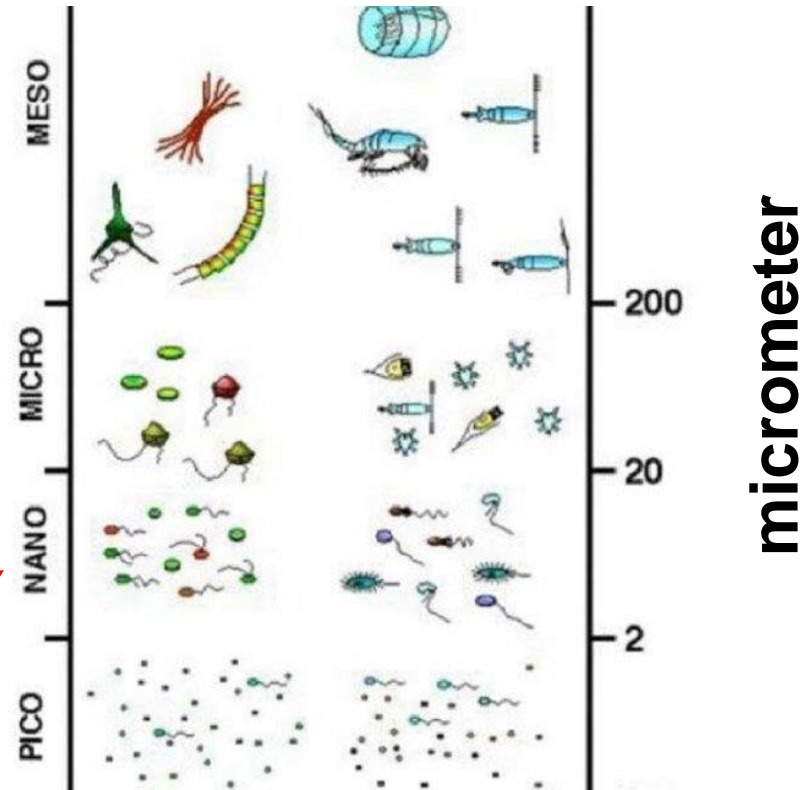
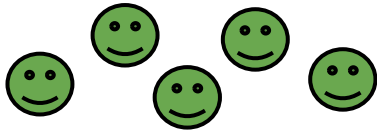
Why is this work needed?

- Pico eukaryotes and pico cyanobacteria have different effects on ecosystems (cyanobacteria such as *Cyanobium* spp. can produce toxins)

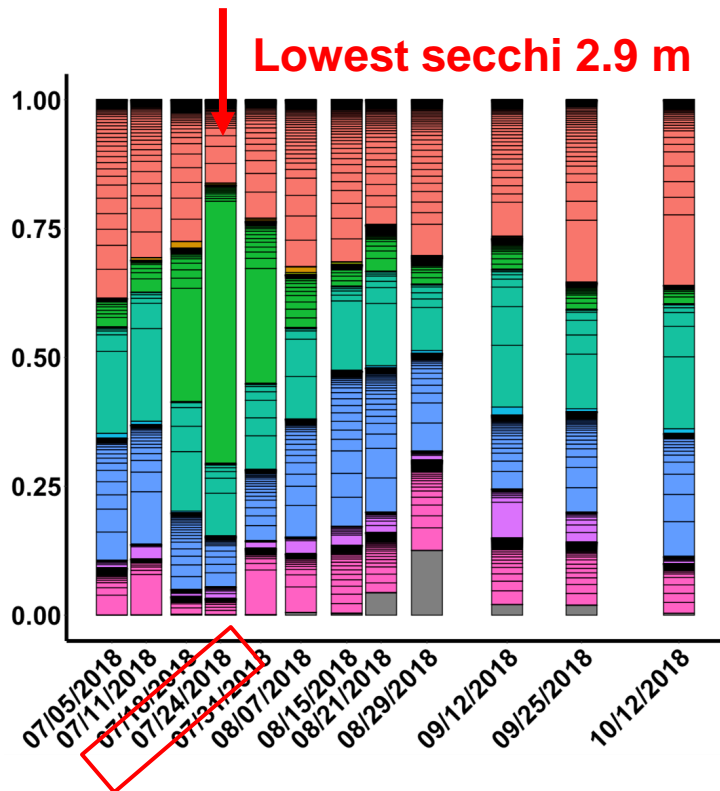
Why use eDNA?

- Unable to identify the taxa responsible for decreasing water transparency using microscopy

L.G.B.



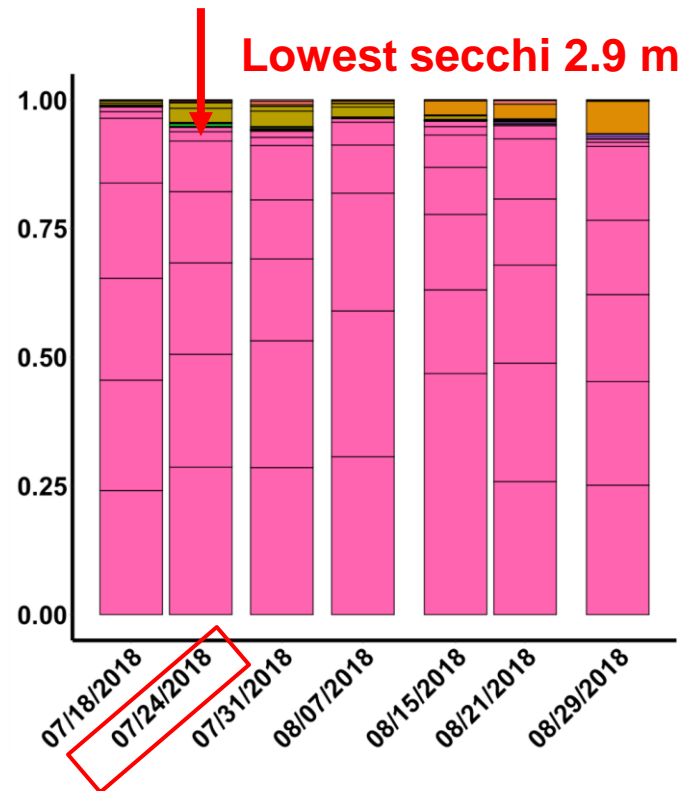
Alcaraz & Calbet (2003)



Super Order Eukaryote (18s)

- Alveolata
- Amoebozoa
- Apusozoa
- Archaeplastida
- Hacrobia
- Hacrobia:nucl
- Opisthokonta
- Rhizaria
- Stramenopiles
- NA

Rhexinema spp.
(formerly known
as *Helicodictyon*)

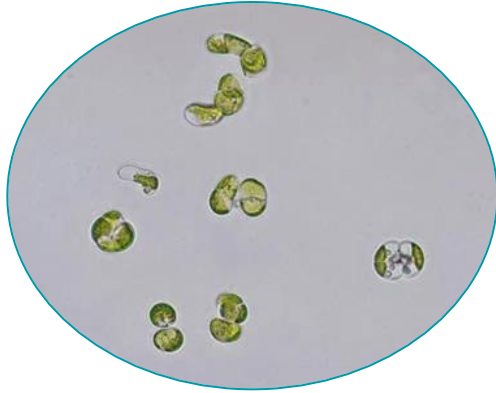


Order Bacteria (16s)

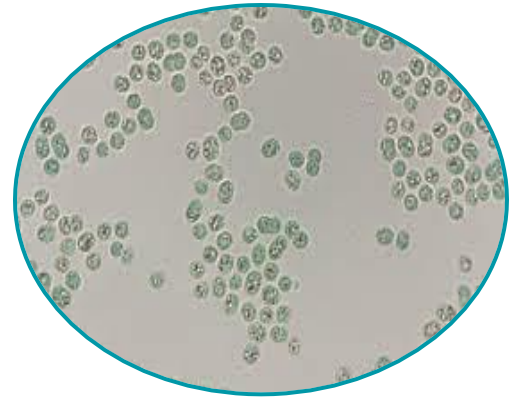
- Actinobacteria
- Chroococcales
- Cyanobacteriales
- Firmicutes
- Hypomicrobiales
- Parcubacteria
- Planctomycetes
- Pseudanabaenales
- Sphingomonadales
- Synechococcales

Cyanobium spp.
(pico-cyanobacteria)

Figures provided by
Robin Sleith



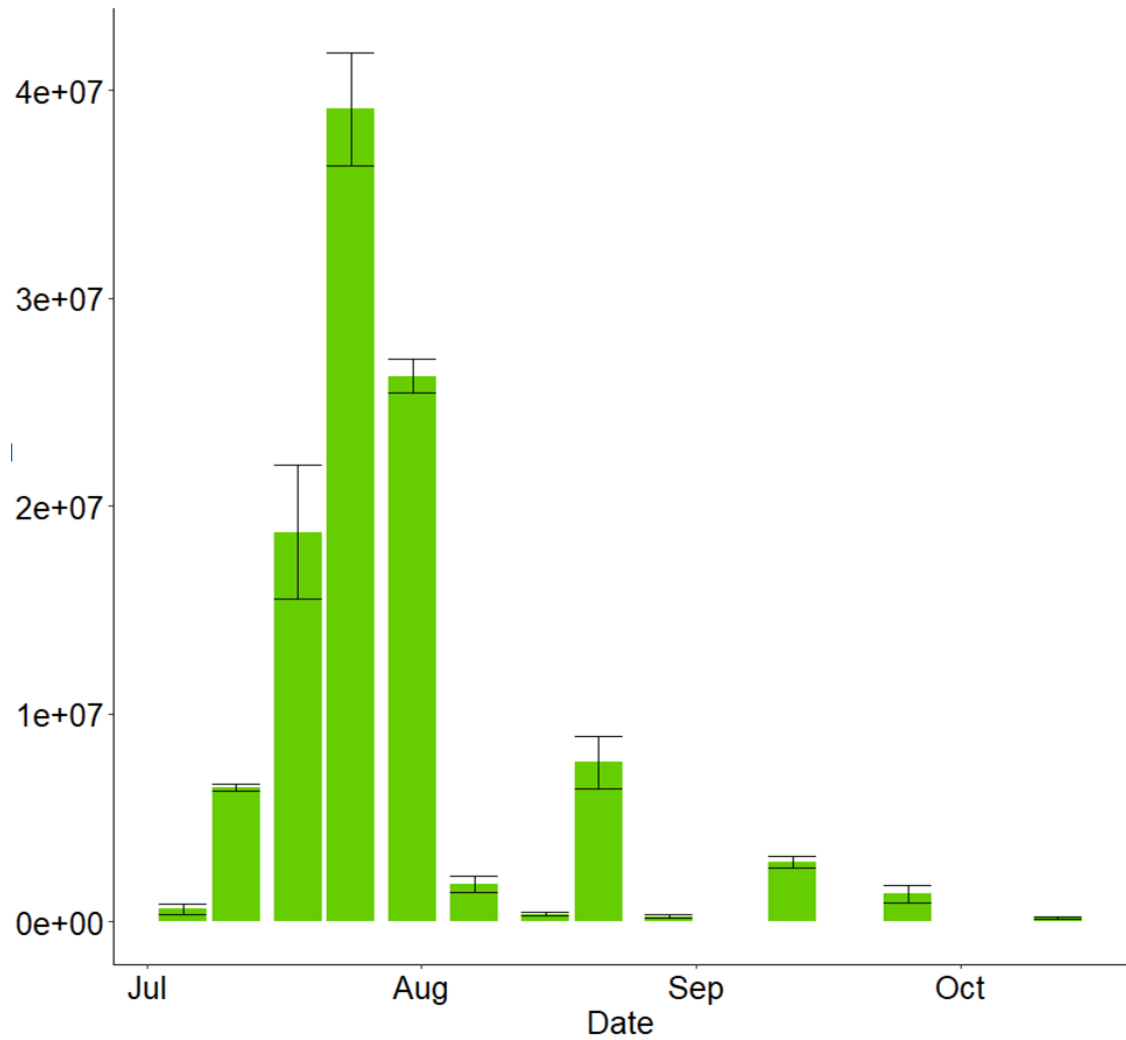
Rhexinema (genus of green algae)
One cell $\leq 10 \mu\text{m}$

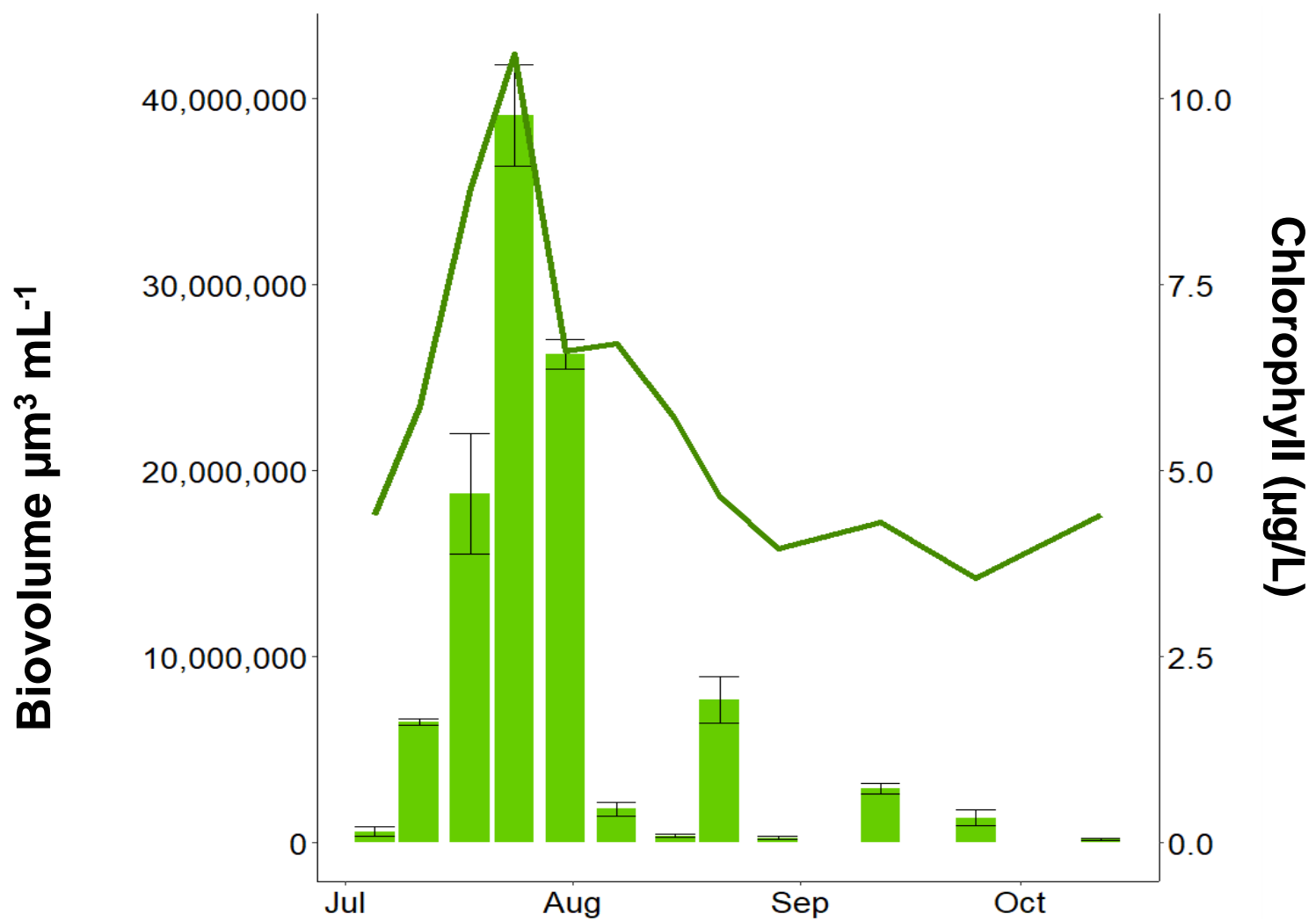


Cyanobium (genus of cyanobacteria)
One cell $\leq 2 \mu\text{m}$

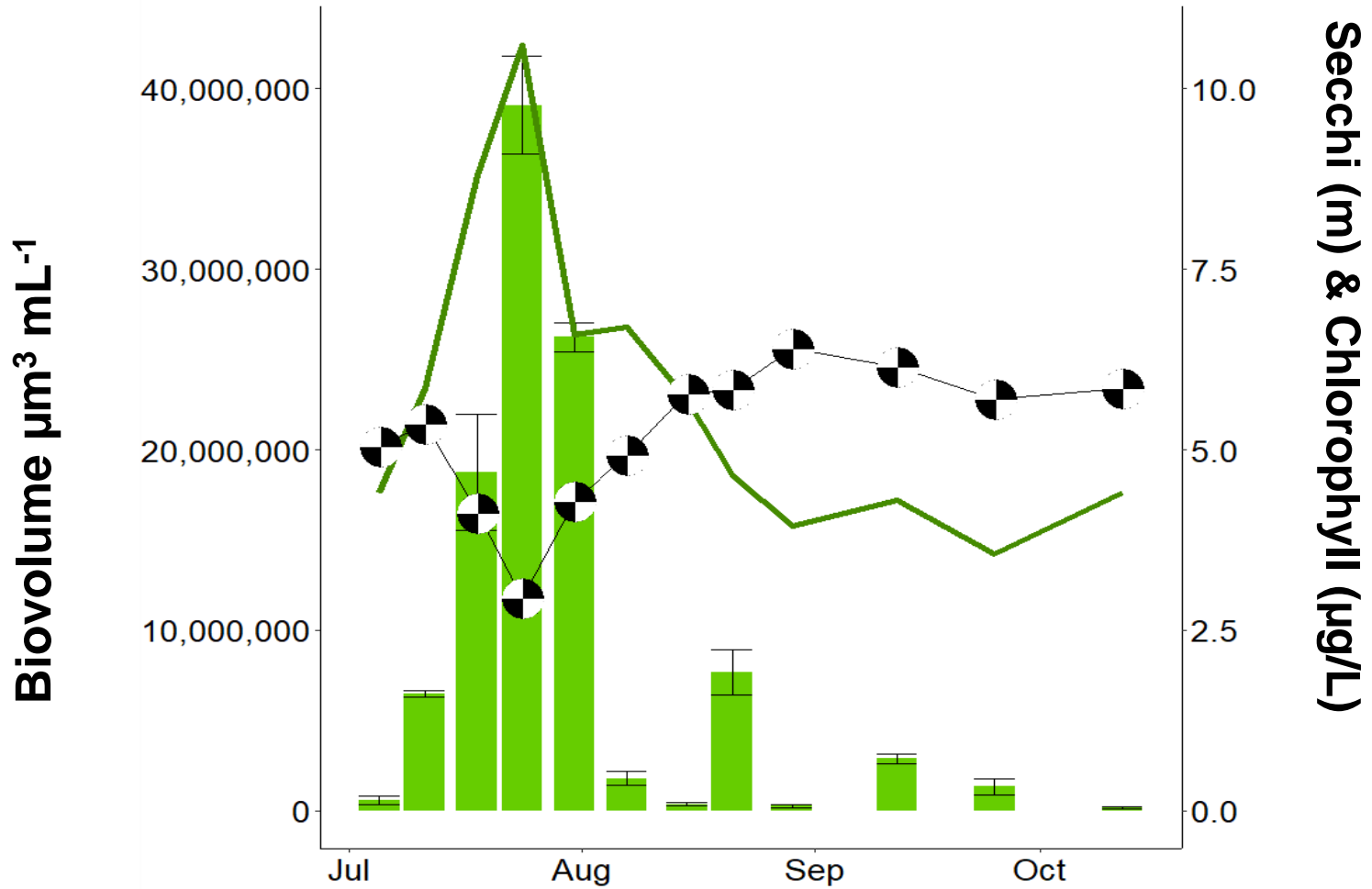
- Design qPCR assays
- Quantify cells volume in epilimnetic water samples

Biovolume $\mu\text{m}^3 \text{mL}^{-1}$





Rhexinema, 2018



2018

Rhexinema (green algae)

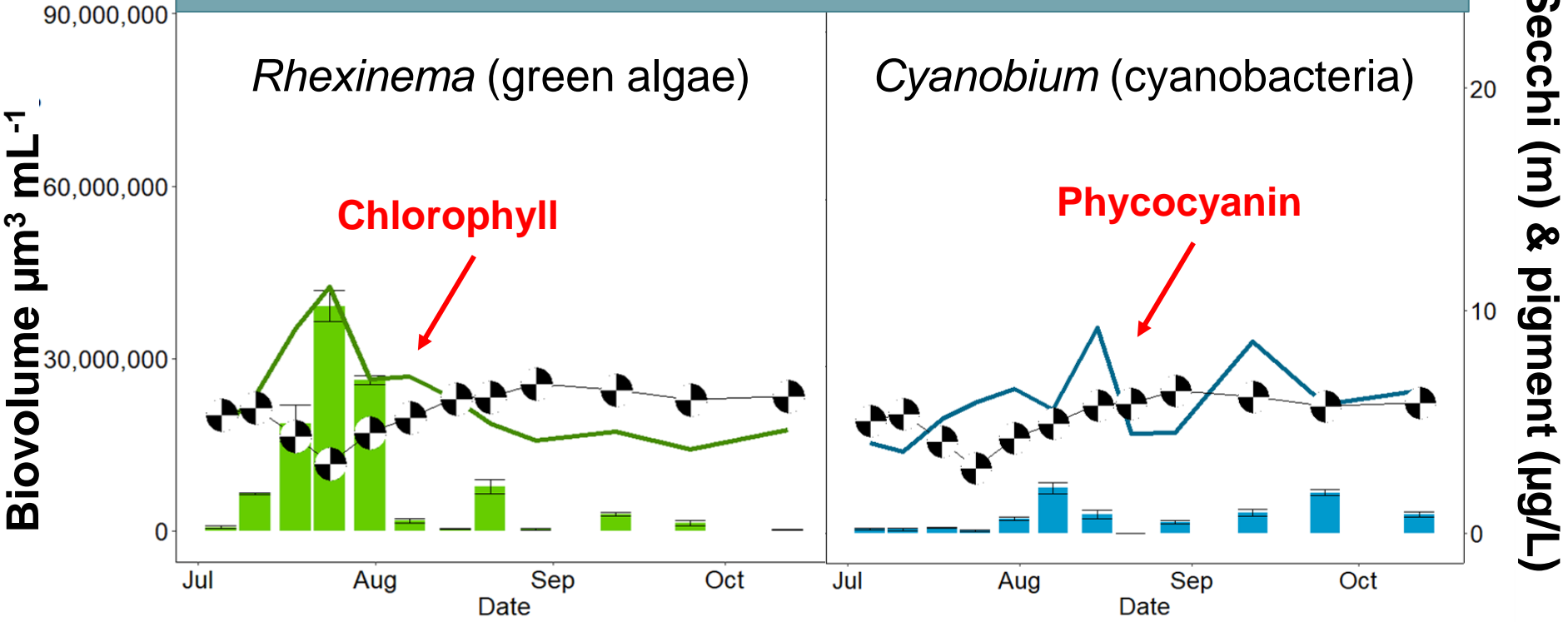
Cyanobium (cyanobacteria)

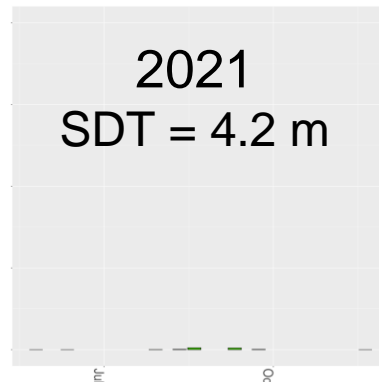
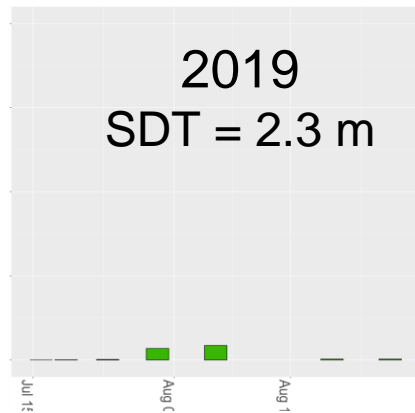
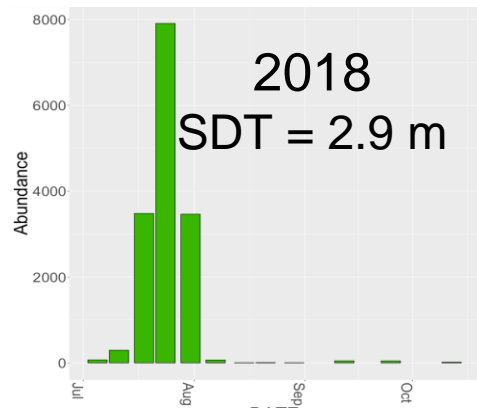
Chlorophyll

Phycocyanin

Biovolume $\mu\text{m}^3 \text{mL}^{-1}$

Secchi (m) & pigment ($\mu\text{g/L}$)

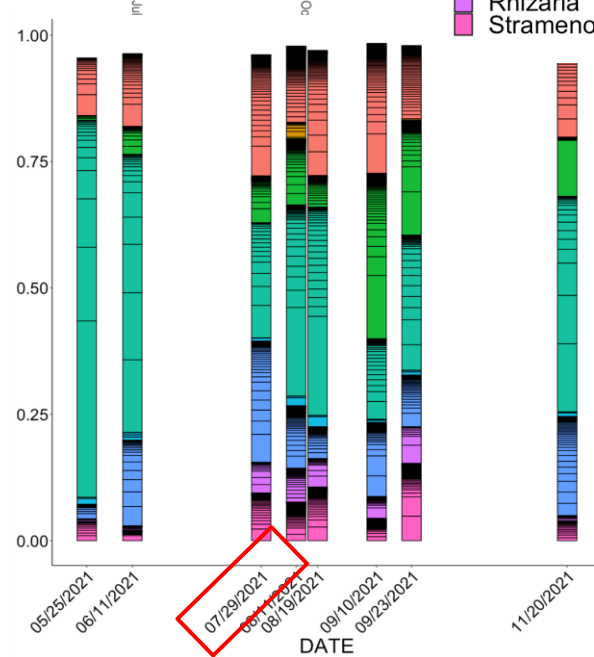
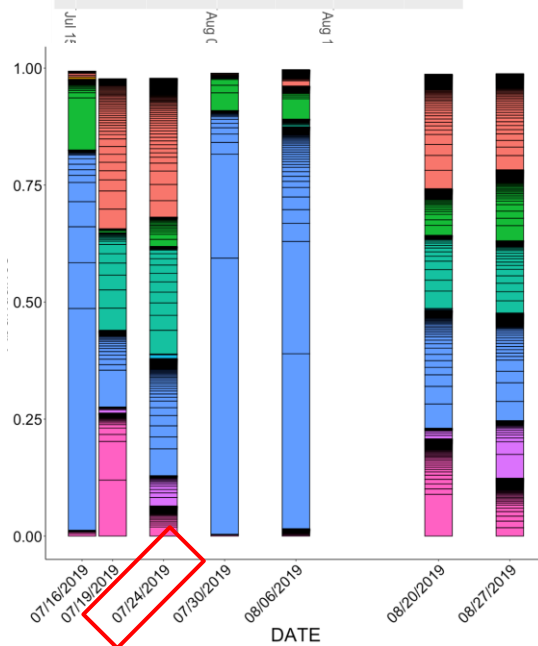
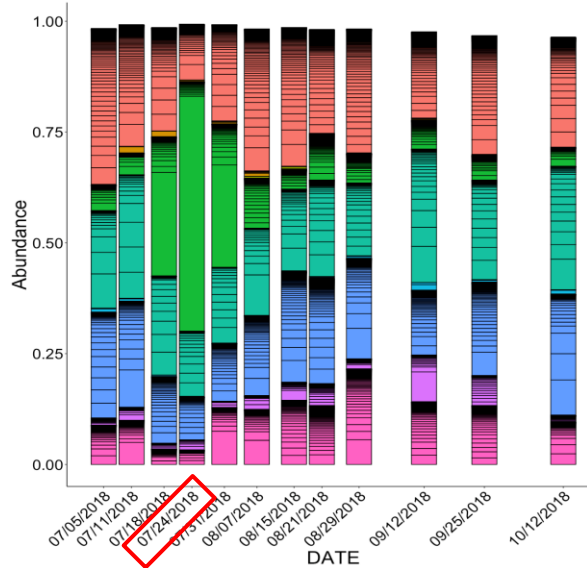




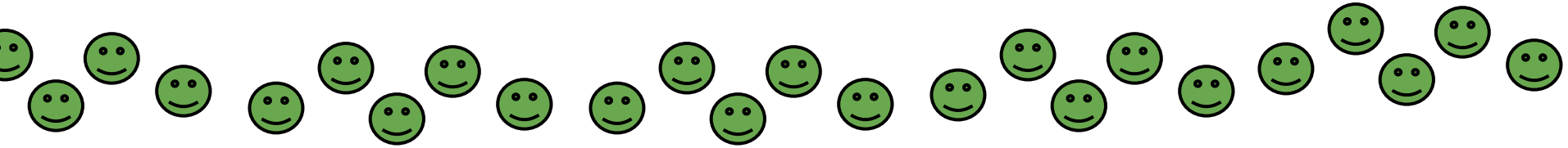
OTU_name
■ ASV_0009

tax.SuperGroup

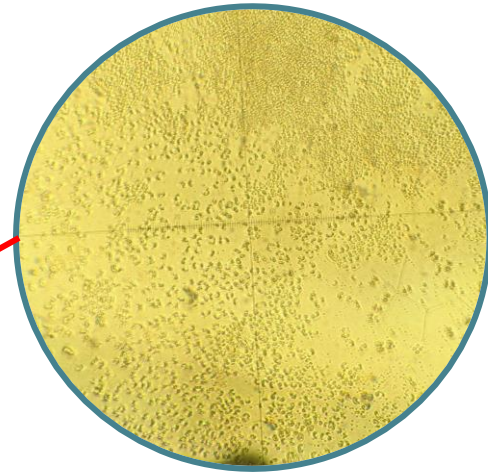
- Alveolata
- Amoebozoa
- Apusozoa
- Archaeplastida
- Hacrobia
- Hacrobia:nucl
- Opisthokonta
- Rhizaria
- Stramenopiles



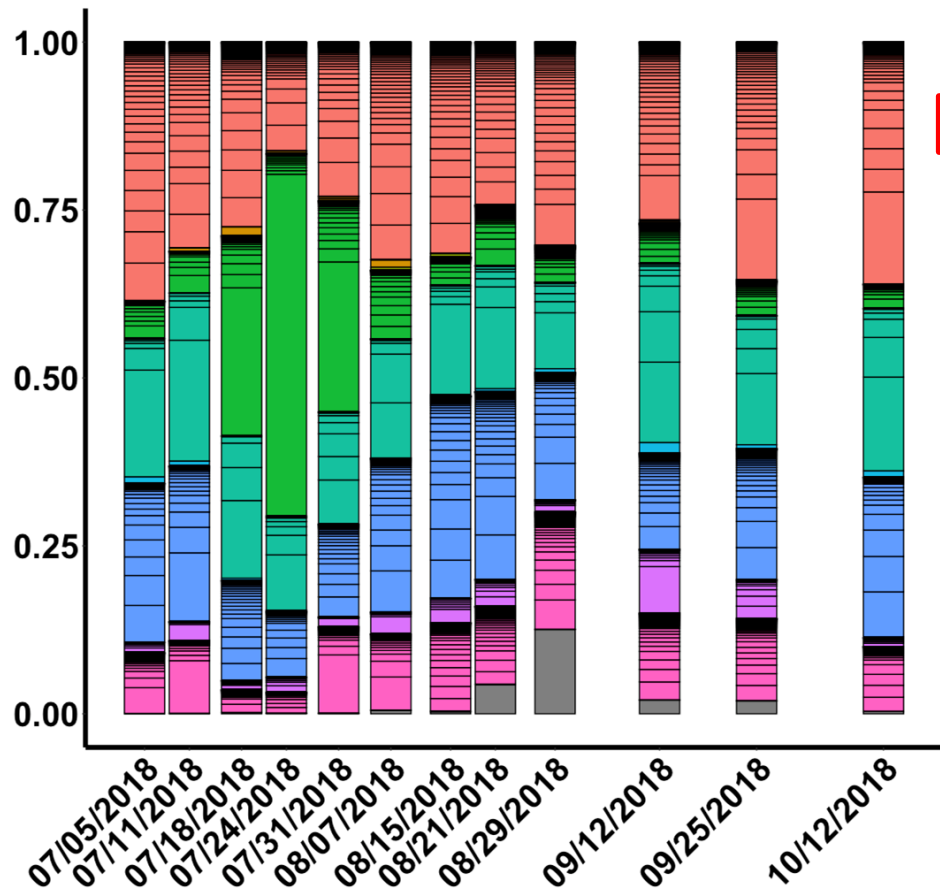
Next Steps



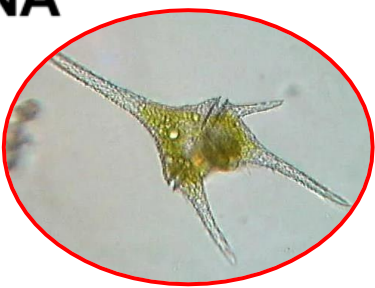
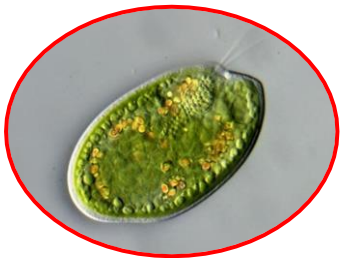
3 meter Secchi
Highland Lake
8/3/2022



- Culture Rhexinema
- Sequence the culture's DNA
- Send culture isolates to a taxonomist



- Alveolata
- Amoebozoa
- Apusozoa
- Archaeplastida
- Hacrobia
- Hacrobia:nucl
- Opisthokonta
- Rhizaria
- Stramenopiles
- NA



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Hannah Braslau



Maddie Rapelyea



Sharon Mann



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