

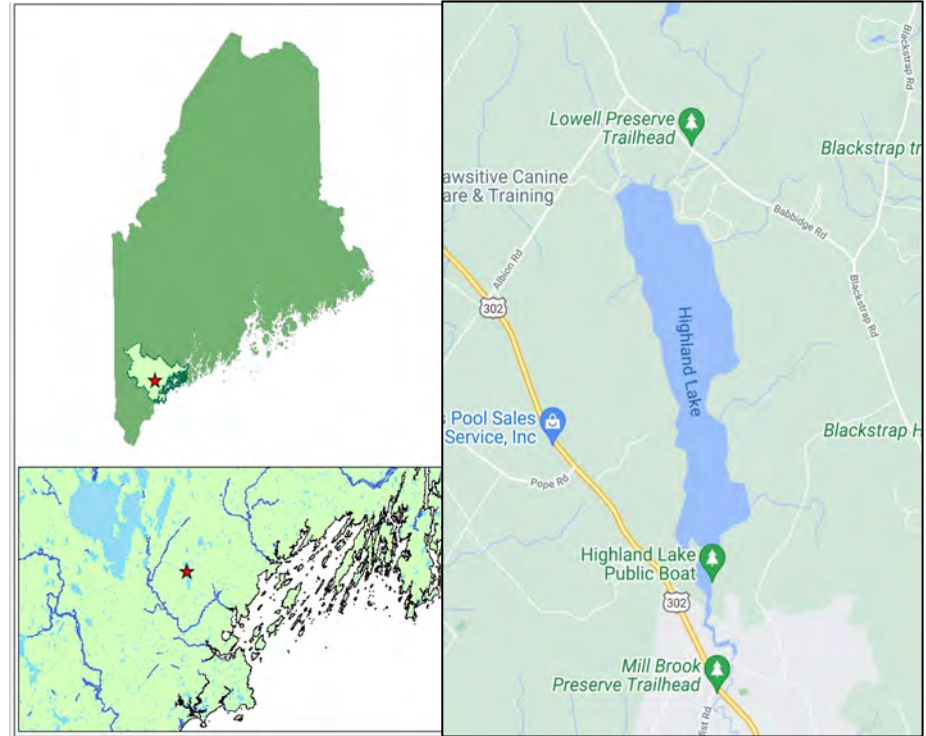
Using eDNA tools to investigate a mysterious algae bloom: An update on Highland (Duck) Lake

Sharon Mann^{1,2}, Karen Wilson¹, Robin Sleith²,
and Peter Countway²

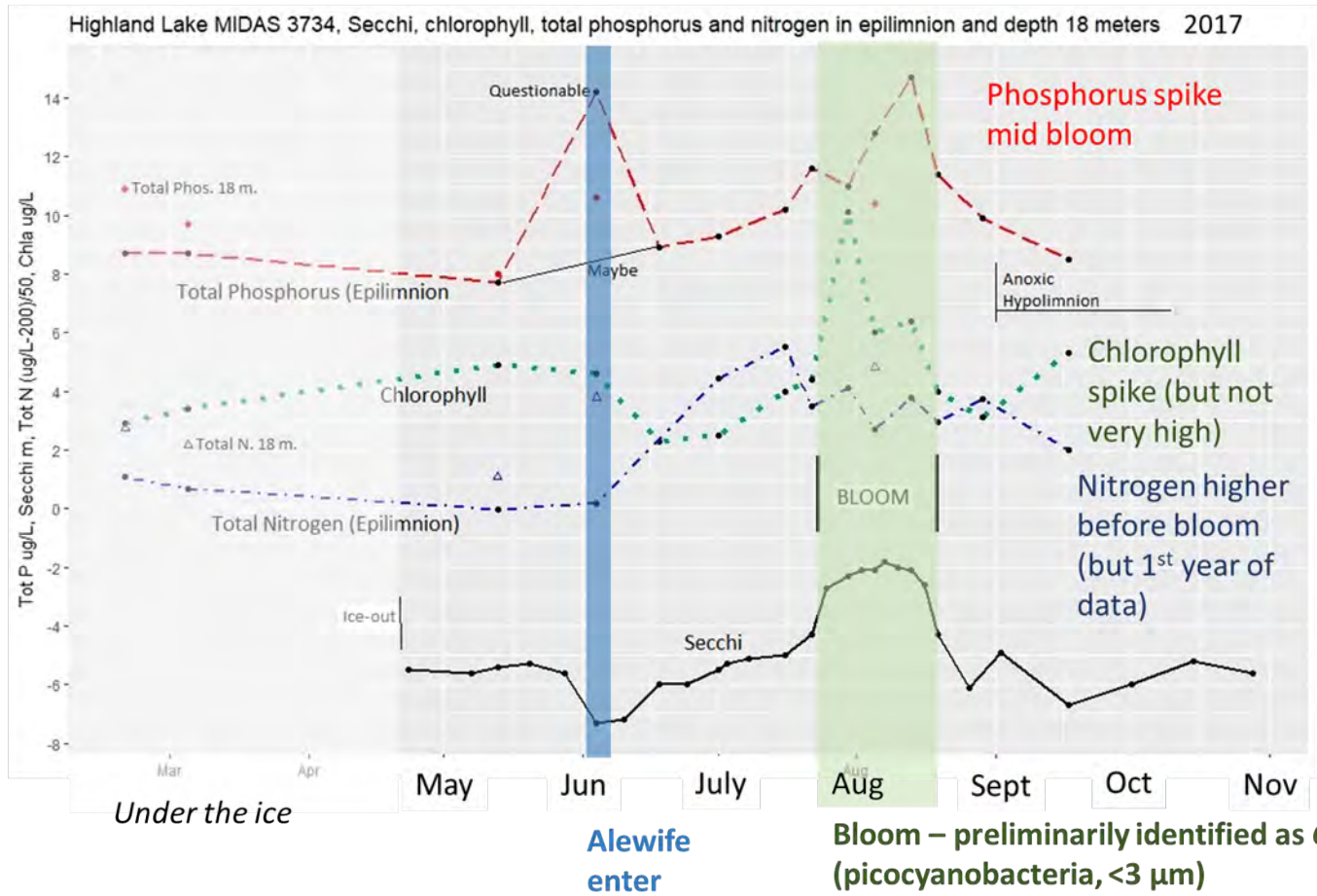
¹University of Southern Maine, ²Bigelow Laboratory for Ocean Sciences



- 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



Bloom years
2014 - 2017

Persisted for up to 3 weeks, late July in to Aug.

Secchi depth < 2.5

Bloom – preliminarily identified as *Cyanobium* (picocyanobacteria, <3 μm)

Source: Highland Lake Association (K. Williams); data from HLA, Maine DEP, USM

Cause(s) of algae bloom?

Excess nutrients?

Unlikely: TP levels lower than those associated with nuisance blooms

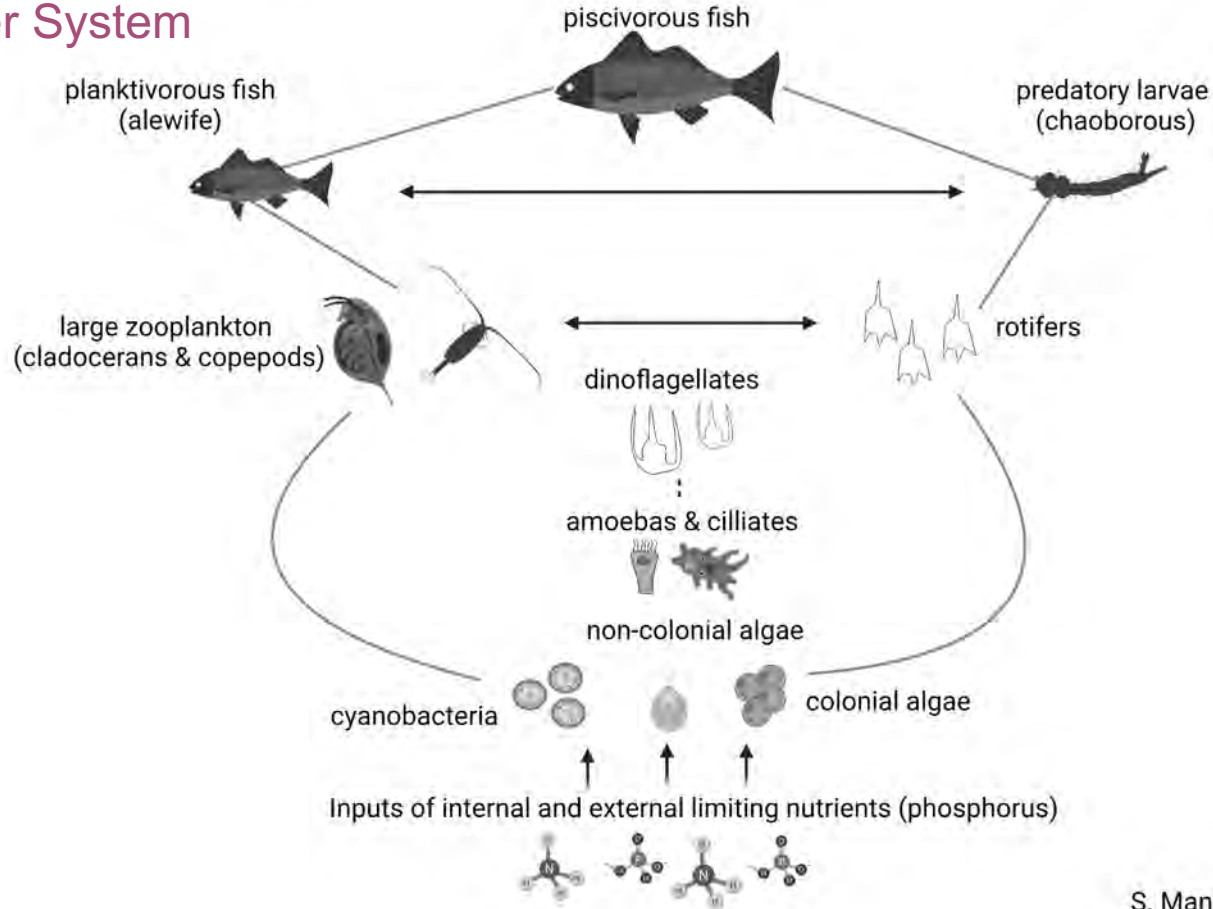
Unusual phytoplankton species?

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

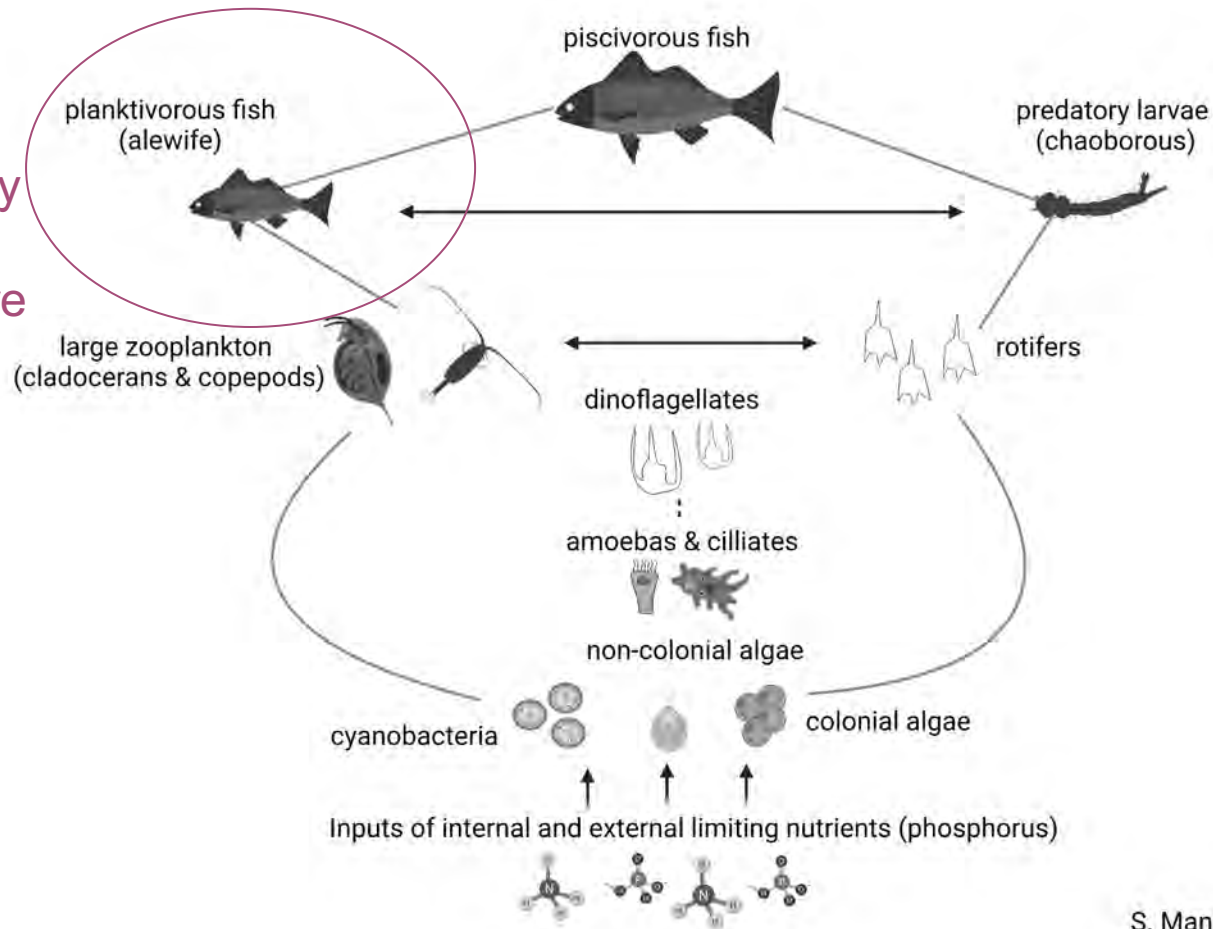
Trophic cascade triggered by consumption of herbivorous zooplankton by alewife?

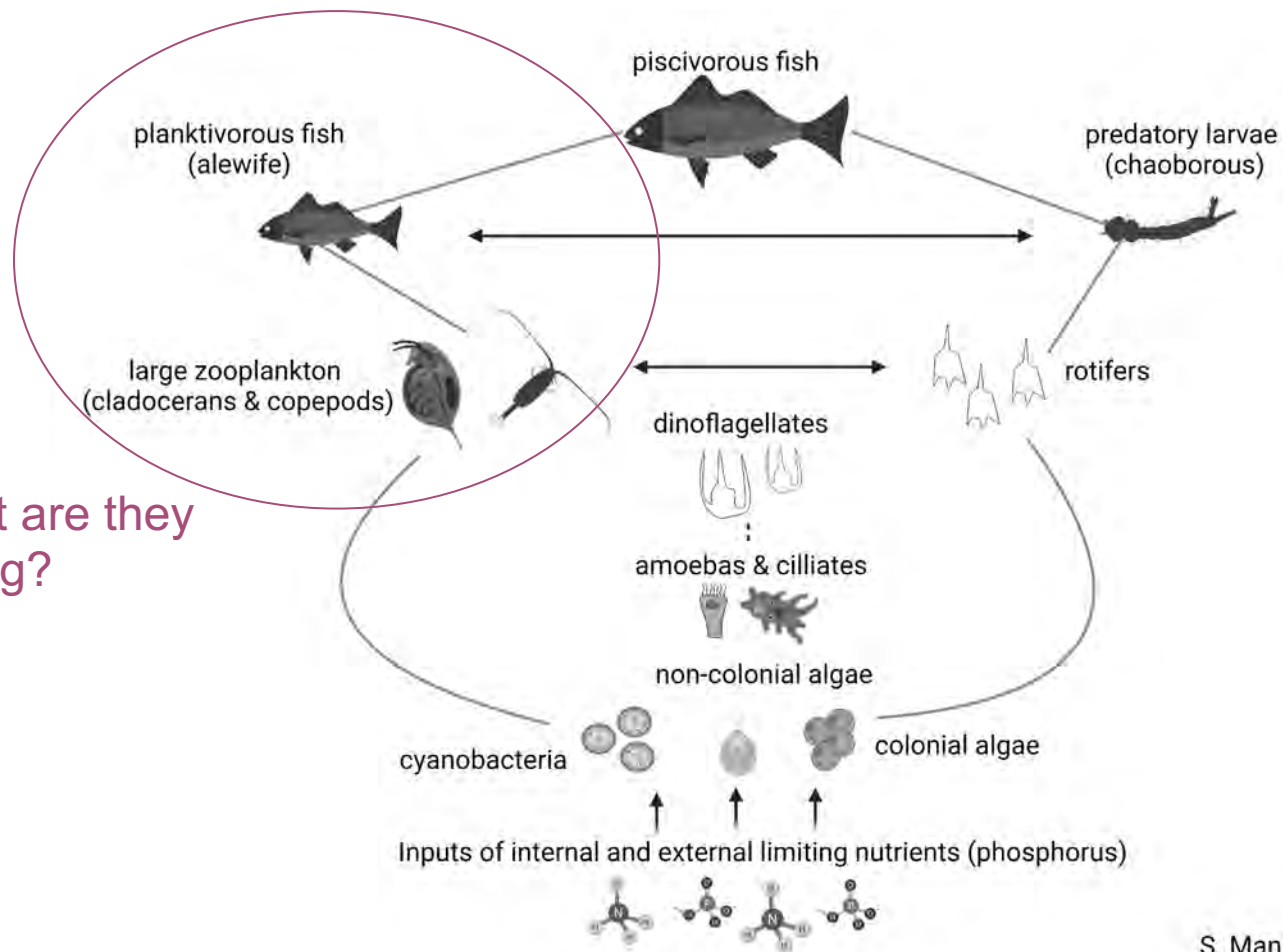
Coming soon!

Freshwater System

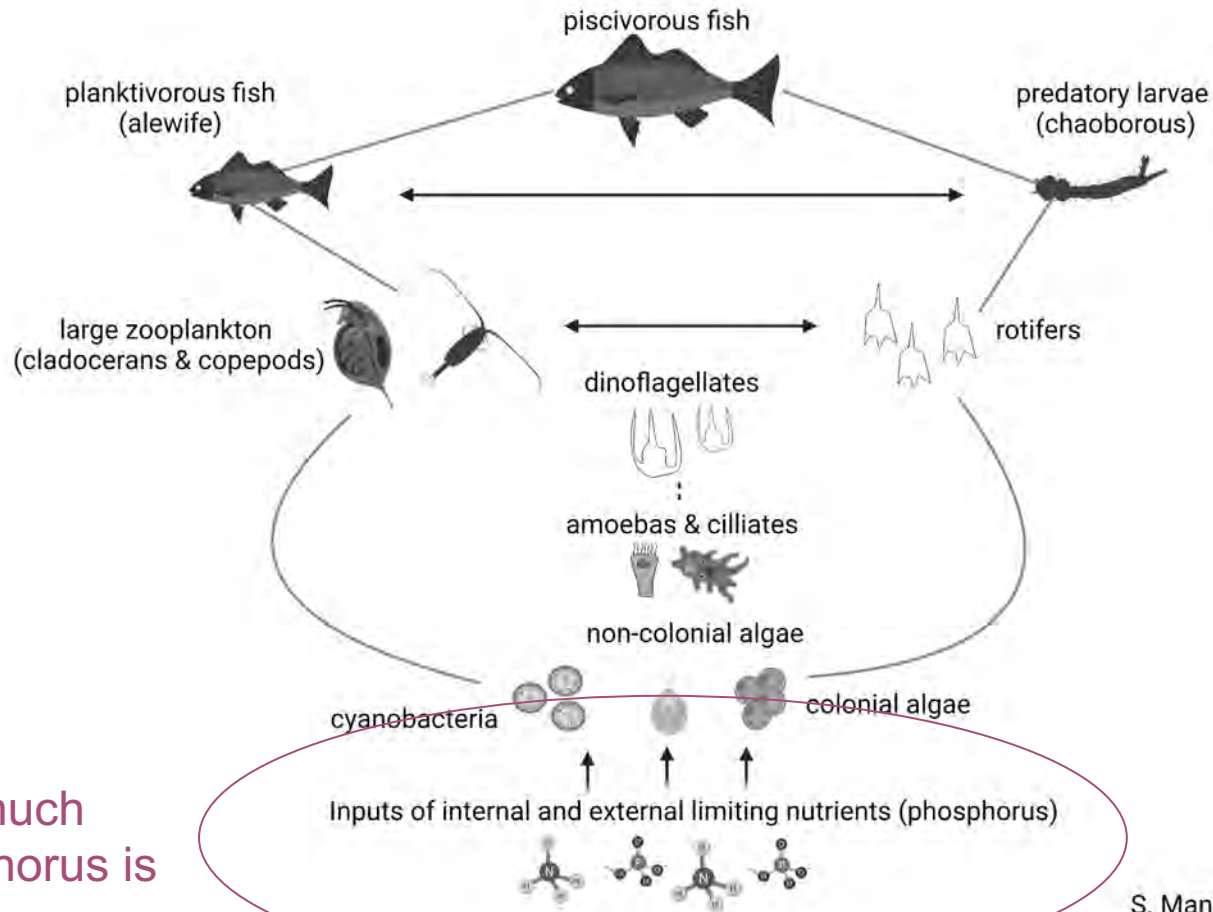


How many juvenile alewife are there?

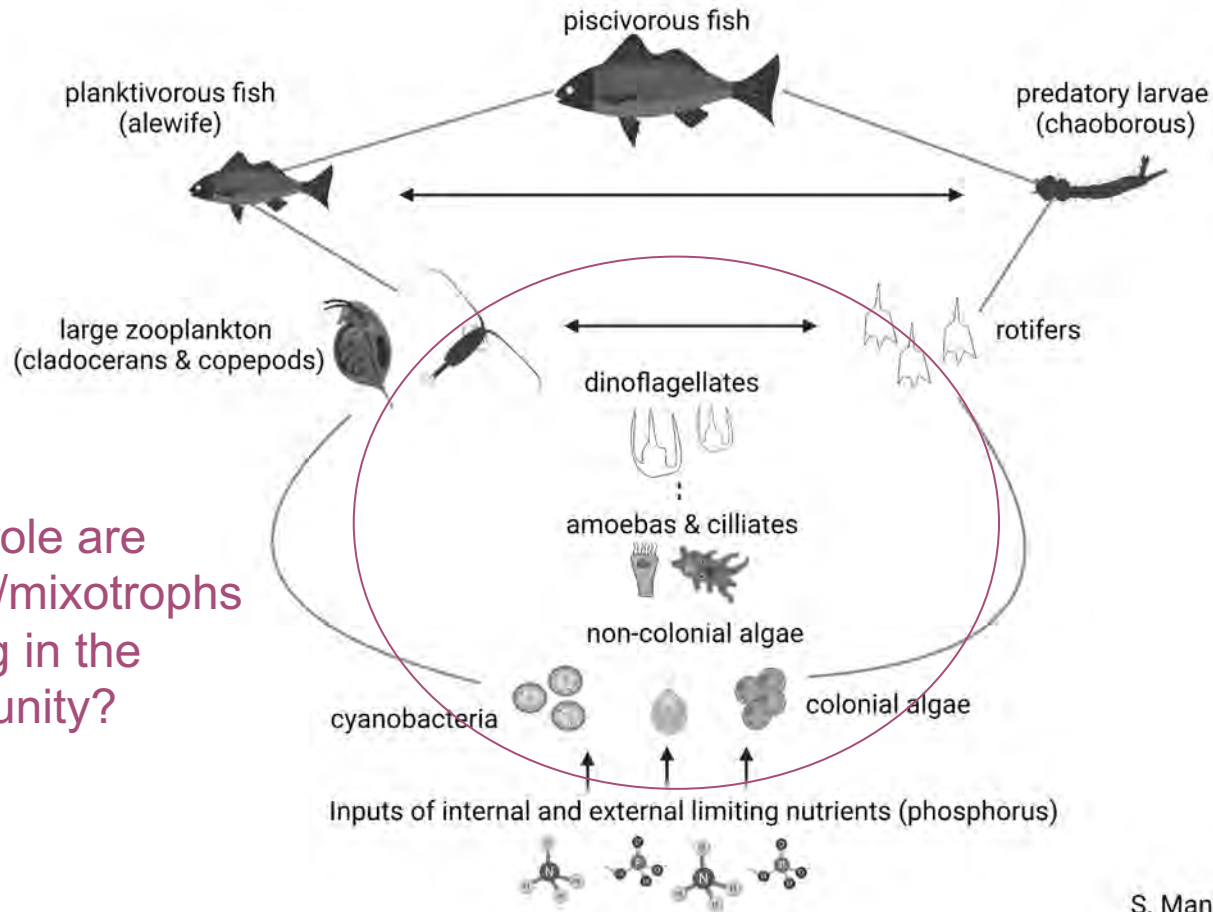




What are they eating?



How much phosphorus is there?



What role are hetero/mixotrophs playing in the community?



Sharon's Dissertation Structure

Part I. Four comparative studies involving eight Maine lakes that differ in base nutrients and anadromous alewife (*Alosa pseudoharengus*) density

Part II. A focused study characterizing the bloom forming taxa in Highland Lake

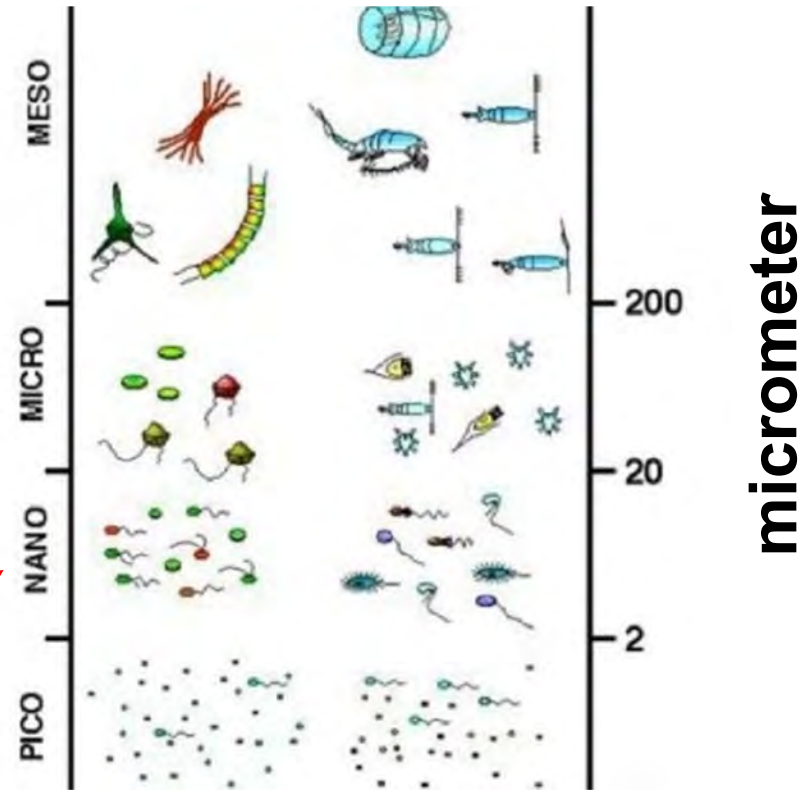
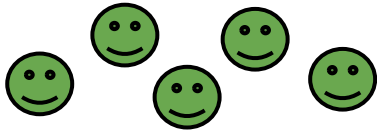
Why is this work needed?

- 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)

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?

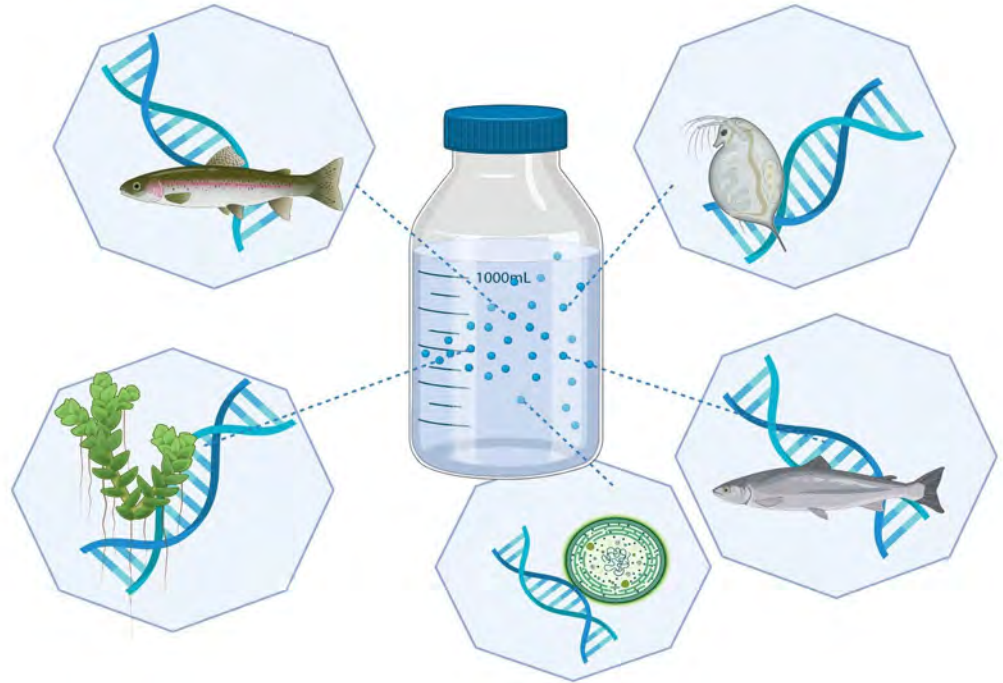
- Cellular decomposition
- Whole shed cells
- Whole microorganisms

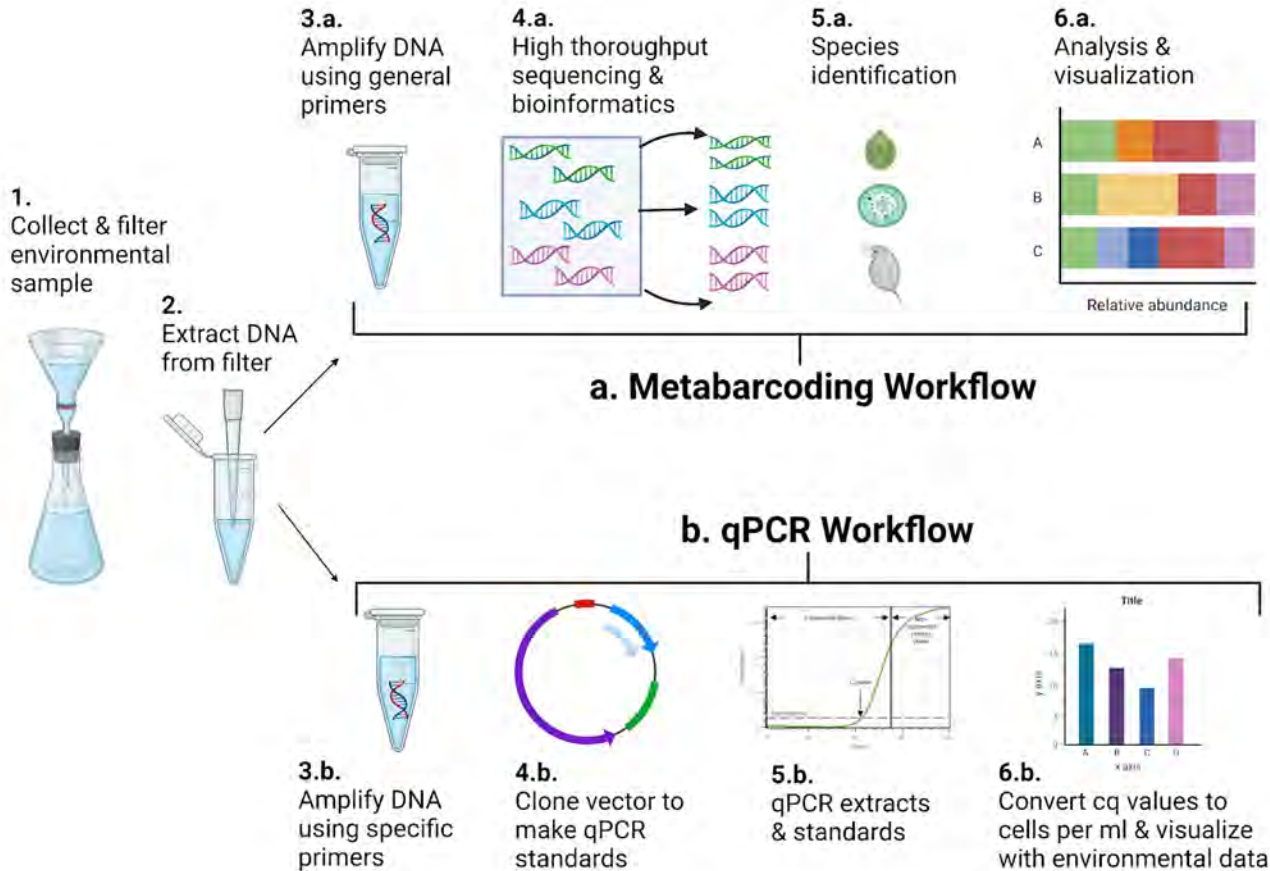
Where is eDNA found?

- Water
- Soils
- Air

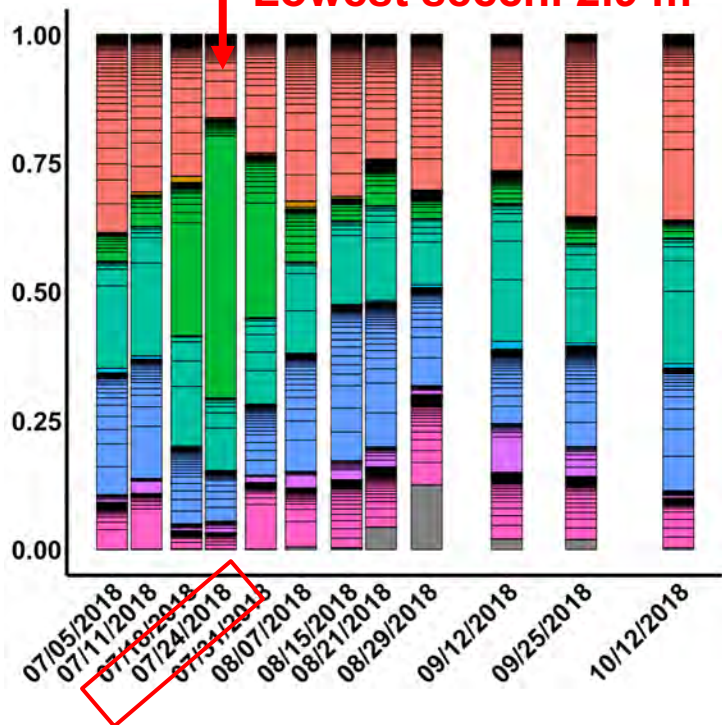
How is eDNA used?

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





Lowest secchi 2.9 m

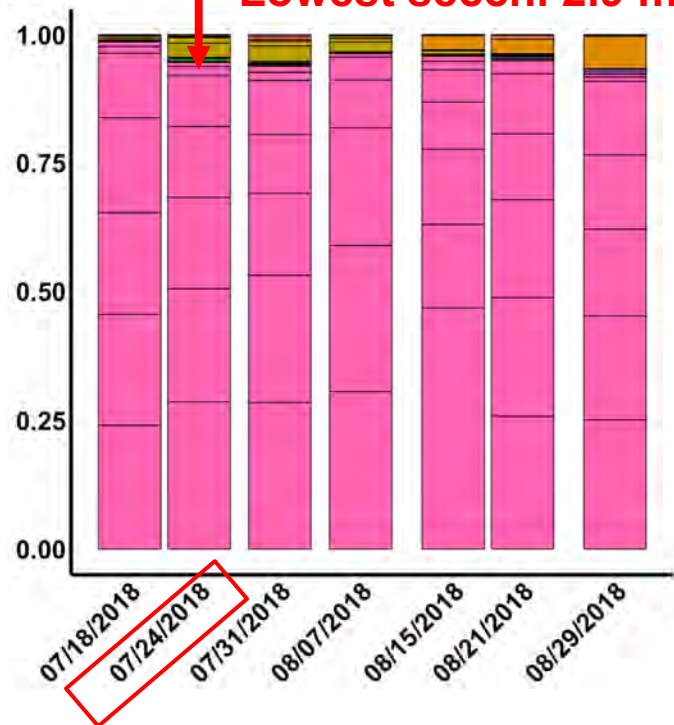


Super Order Eukaryote (18s)

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

Rhexinema spp.
(formerly known
as *Helicodictyon*)

Lowest secchi 2.9 m

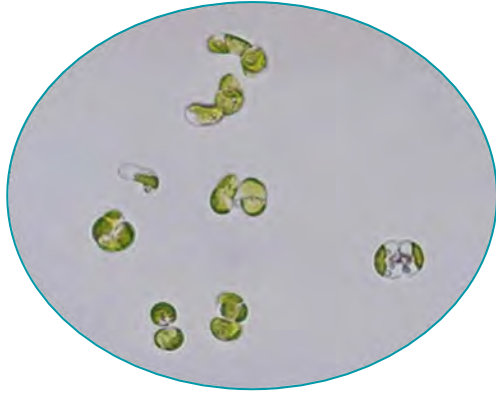


Order Bacteria (16s)

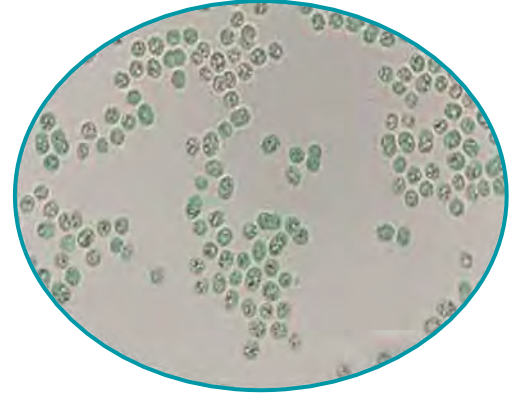
- Actinobacteria
- Chroococcales
- Cyanobacteriales
- Firmicutes
- Hypohomicrobiales
- Parcubacteria
- Planctomycetes
- Pseudanabaenales
- Sphingomonadales
- Synechococcales

Cyanobium spp.
(pico-cyanobacteria)

Figures provided by
Robin Sleith



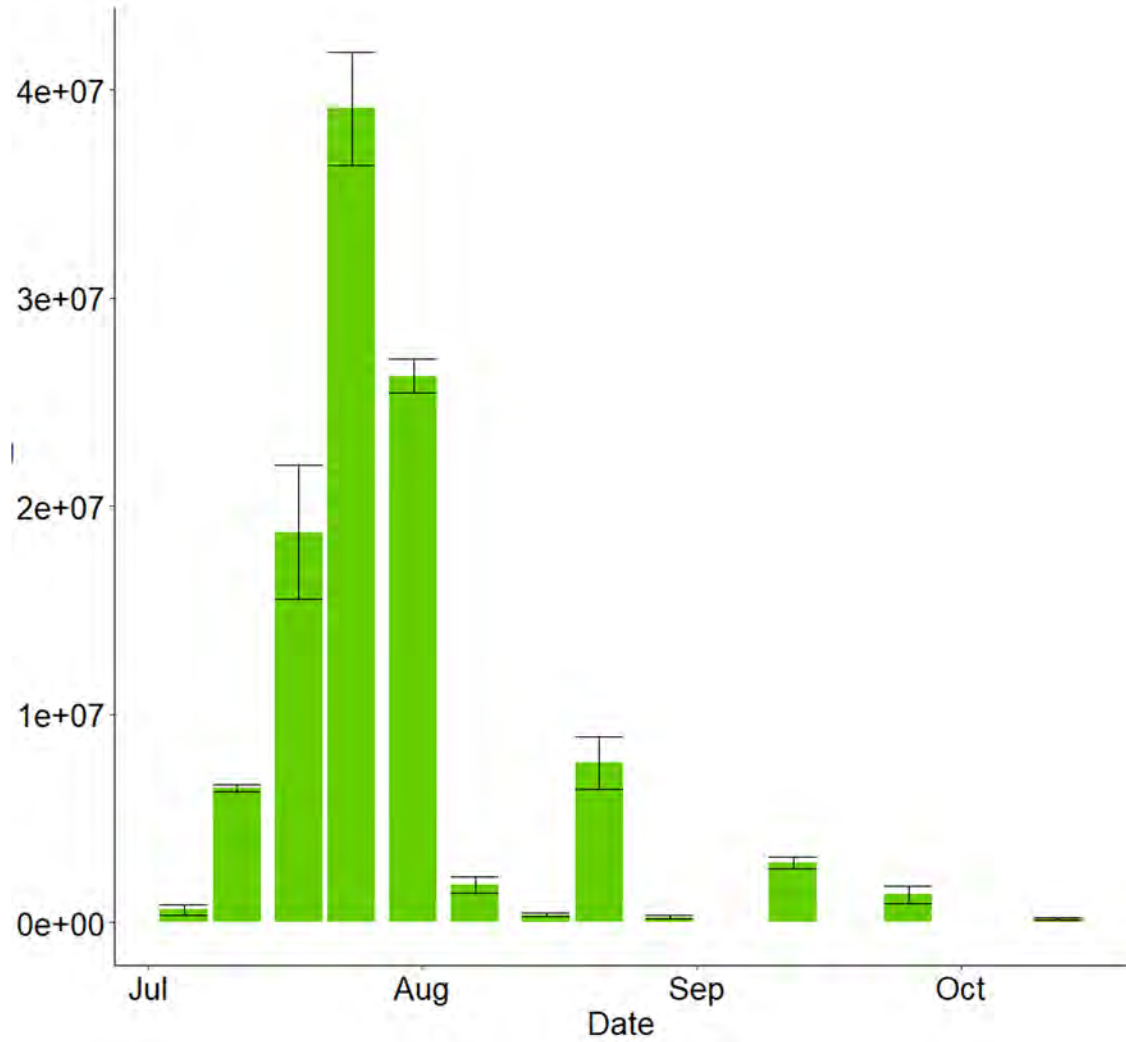
Rhexinema (genus of green algae)
One cell $\leq 10 \mu m$



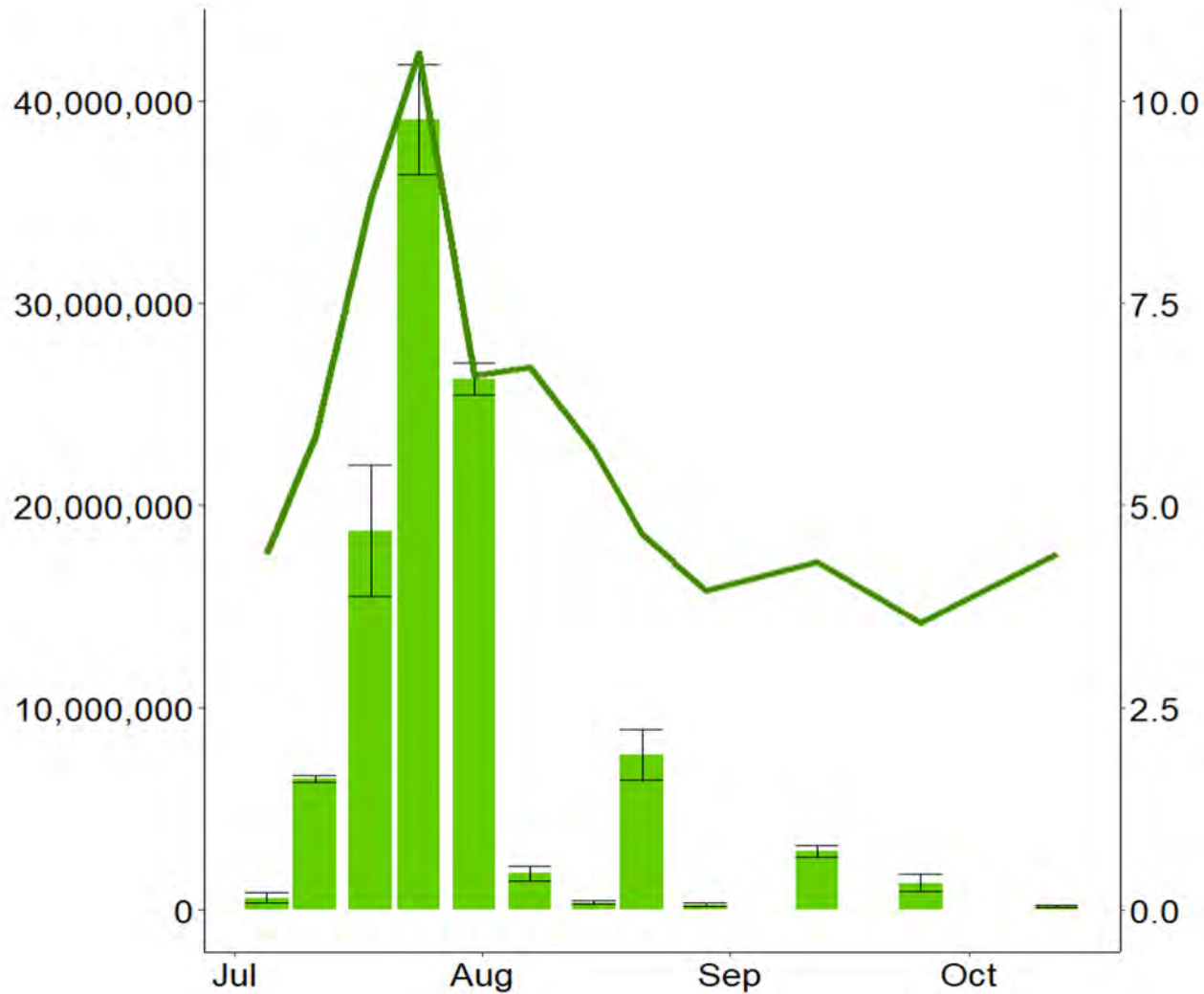
Cyanobium (genus of cyanobacteria)
One cell $\leq 2 \mu m$

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

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



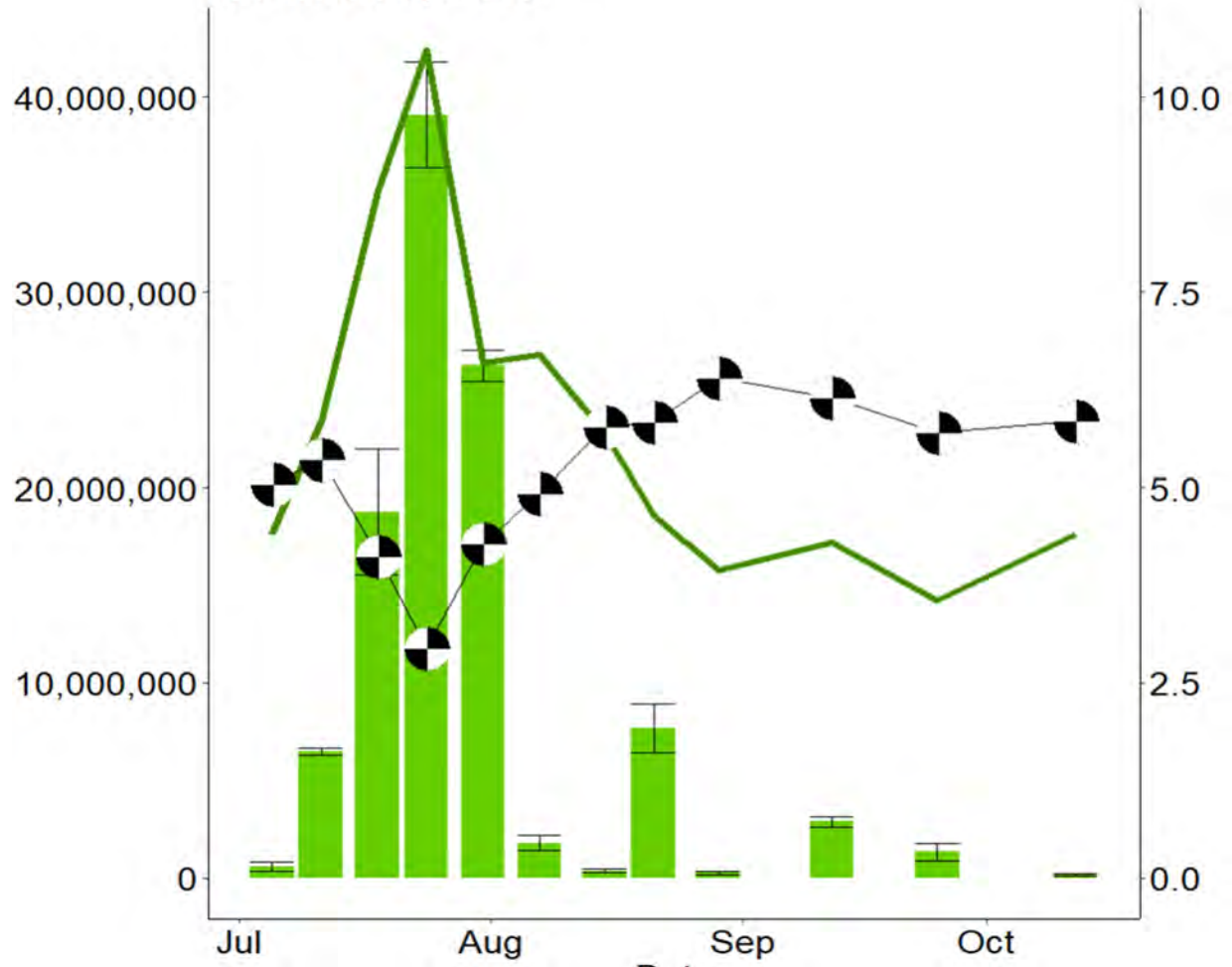
Biovolume $\mu\text{m}^3 \text{mL}^{-1}$



Chlorophyll II ($\mu\text{g/L}$)

Rhexinema, 2018

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



Secchi (m) & Chlorophyll (μg/L)

2018

Rhexinema (green algae)

Cyanobium (cyanobacteria)

Chlorophyll

Phycocyanin

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

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

90,000,000

60,000,000

30,000,000

0

Jul

Aug

Date

Sep

Oct

Jul

Aug

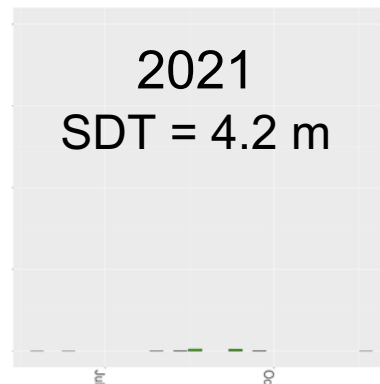
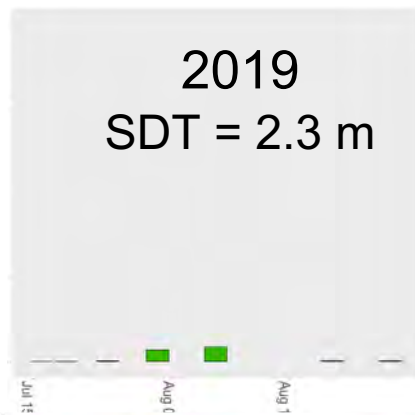
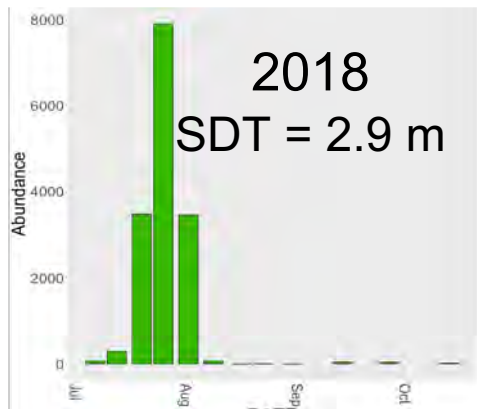
Date

Oct

20

10

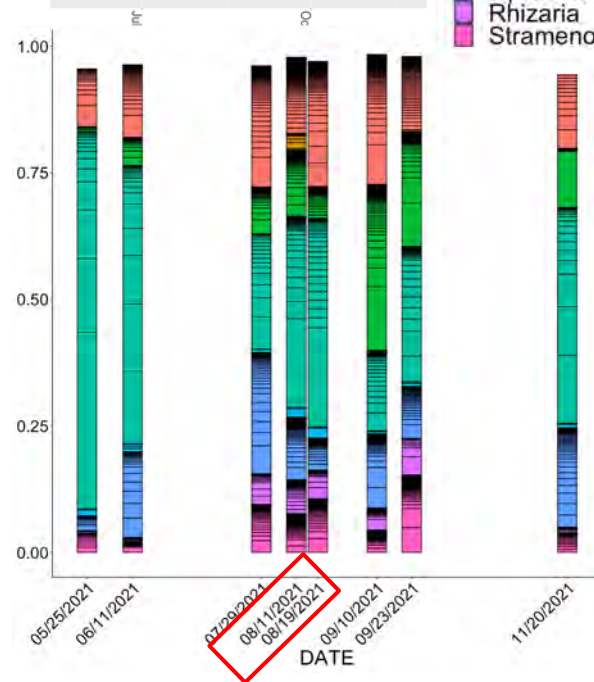
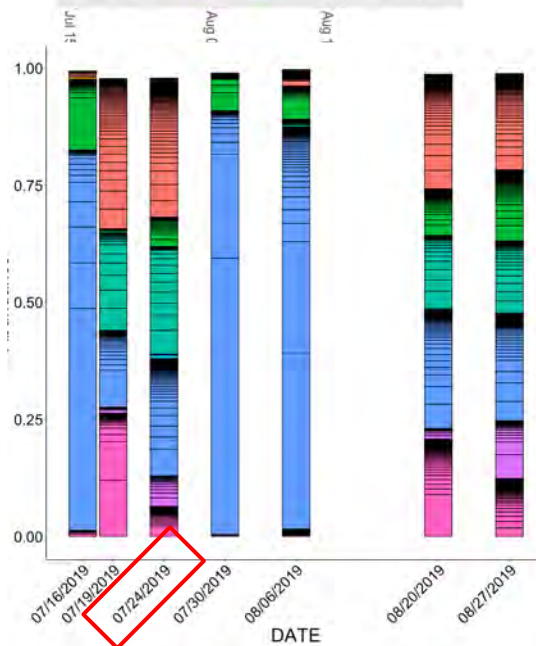
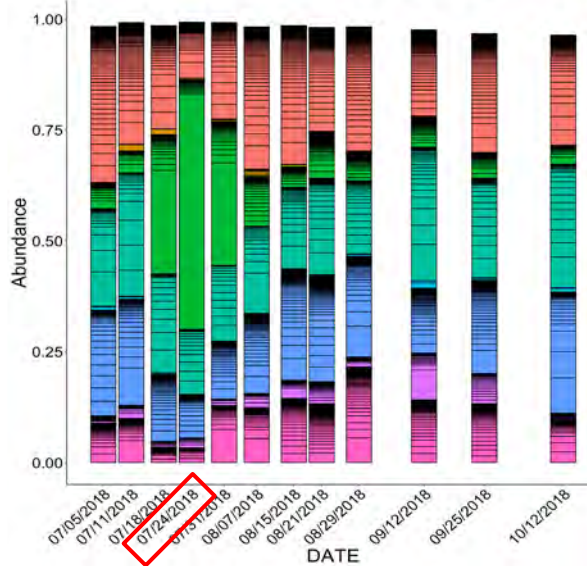
0



OTU_name
ASV_0009

tax.SuperGroup

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

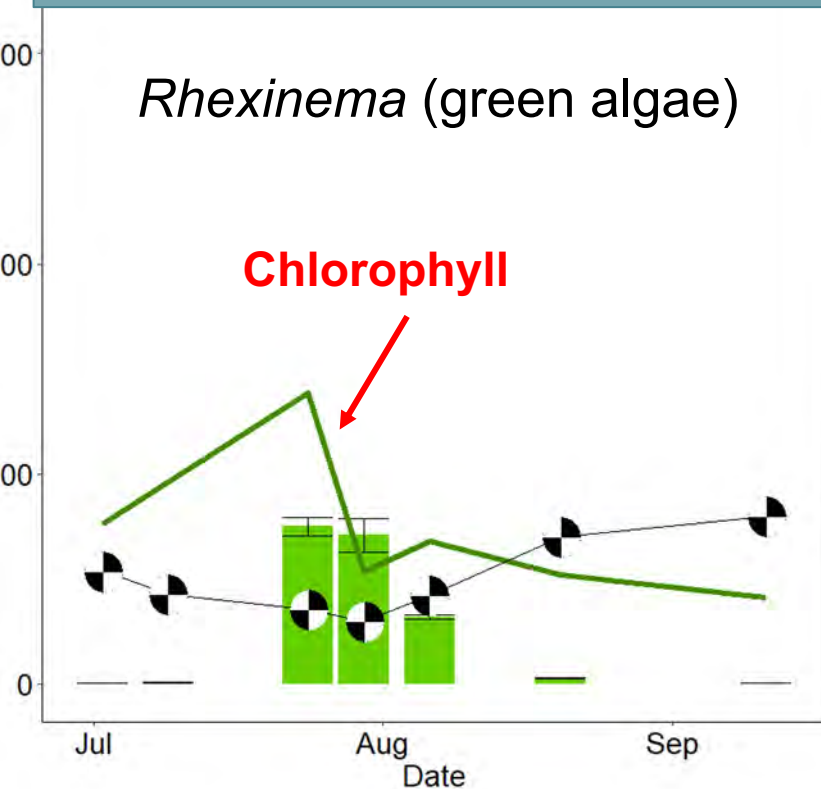


2019

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

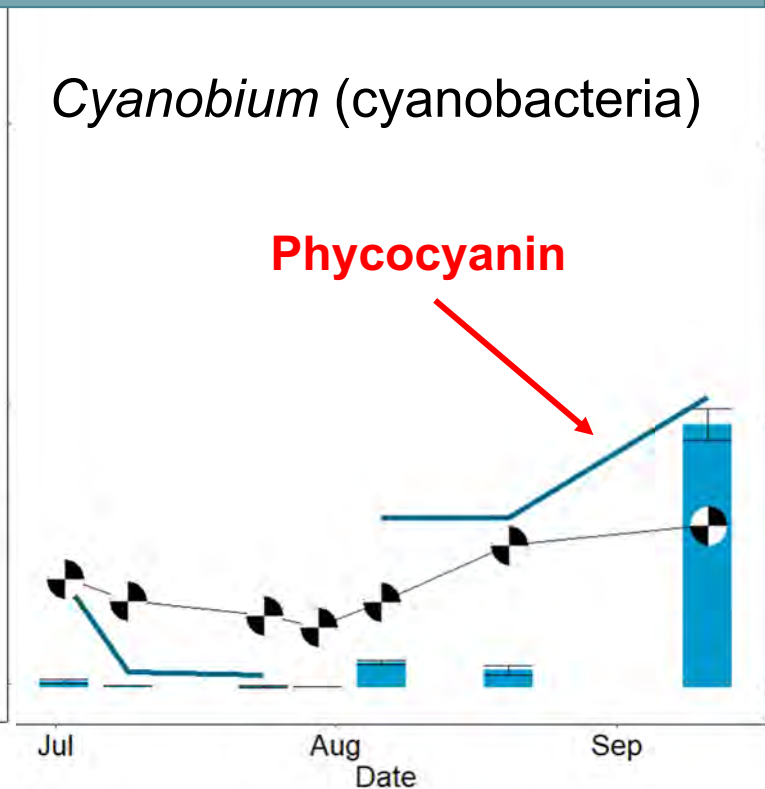
Rhexinema (green algae)

Chlorophyll



Cyanobium (cyanobacteria)

Phycocyanin



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

20

10

0

2021

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

Rhexinema (green algae)

Chlorophyll

90,000,000
60,000,000
30,000,000
0

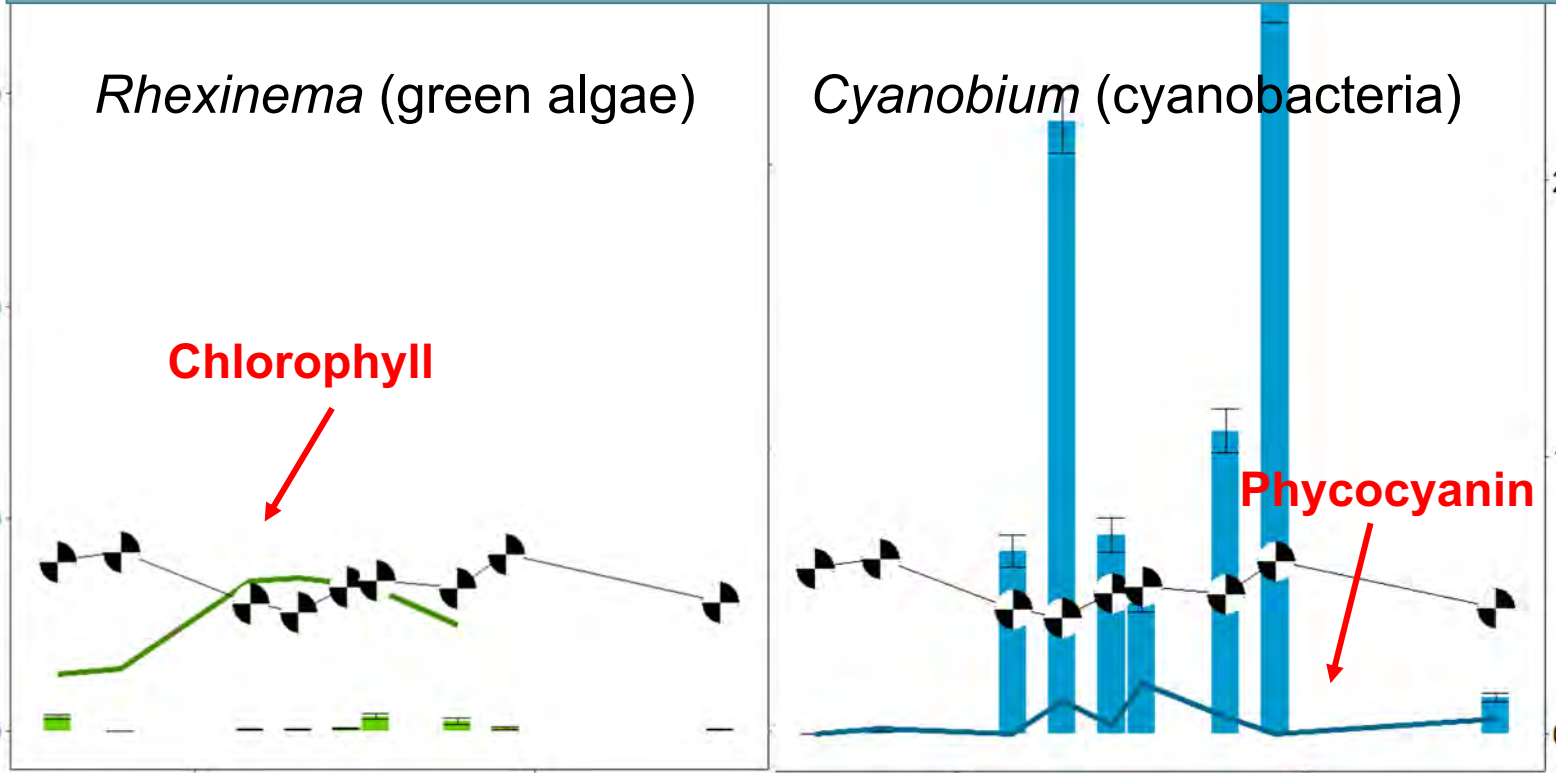
Jul Date Oct

Cyanobium (cyanobacteria)

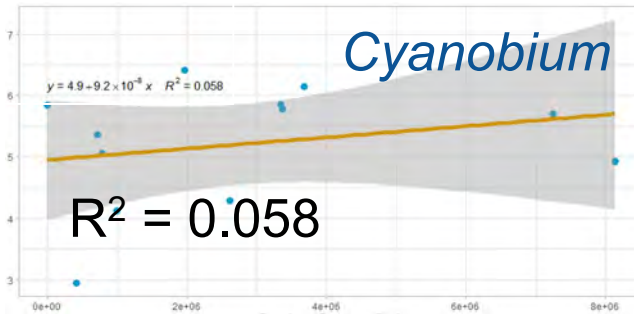
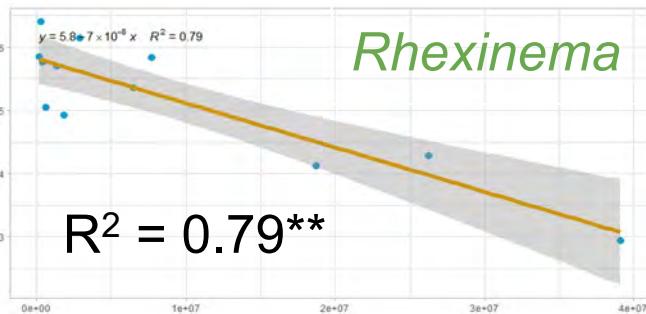
Phycocyanin

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

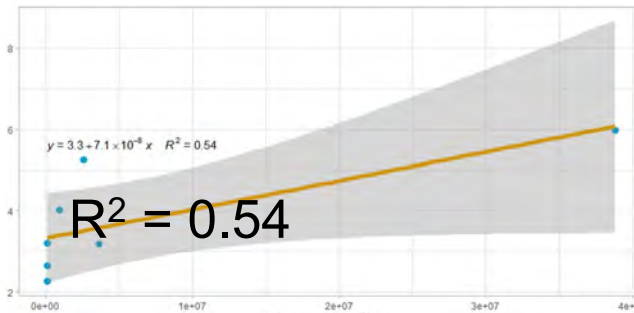
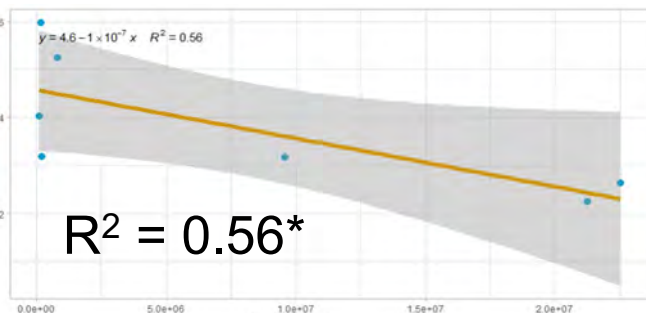
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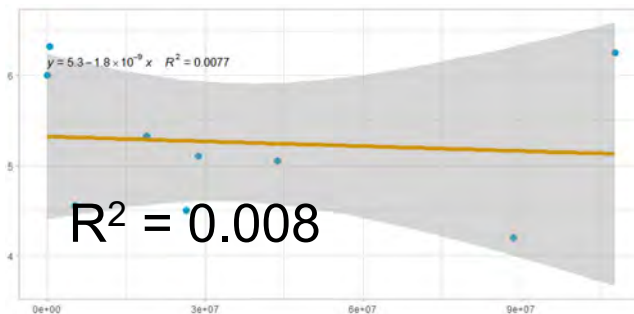
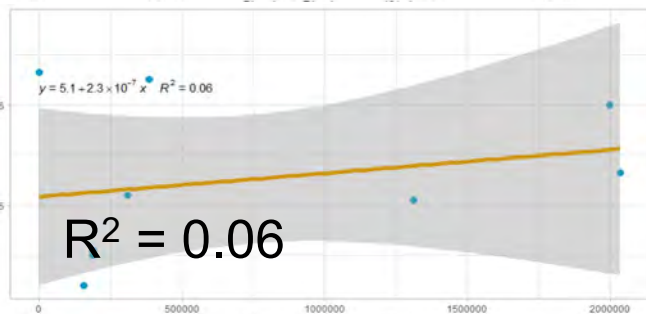
Secchi (m)



2018



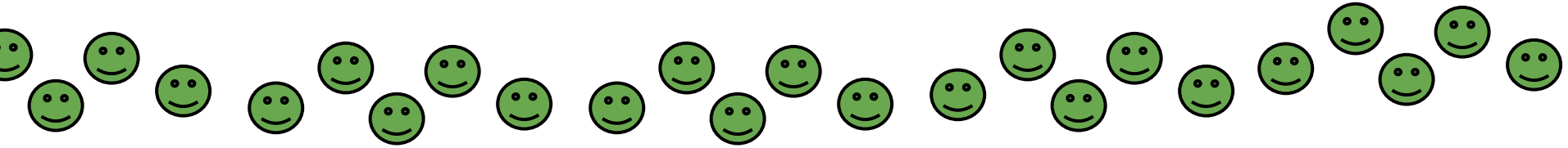
2019



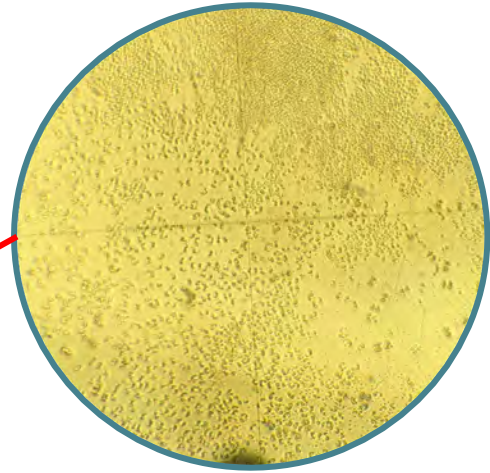
2021

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

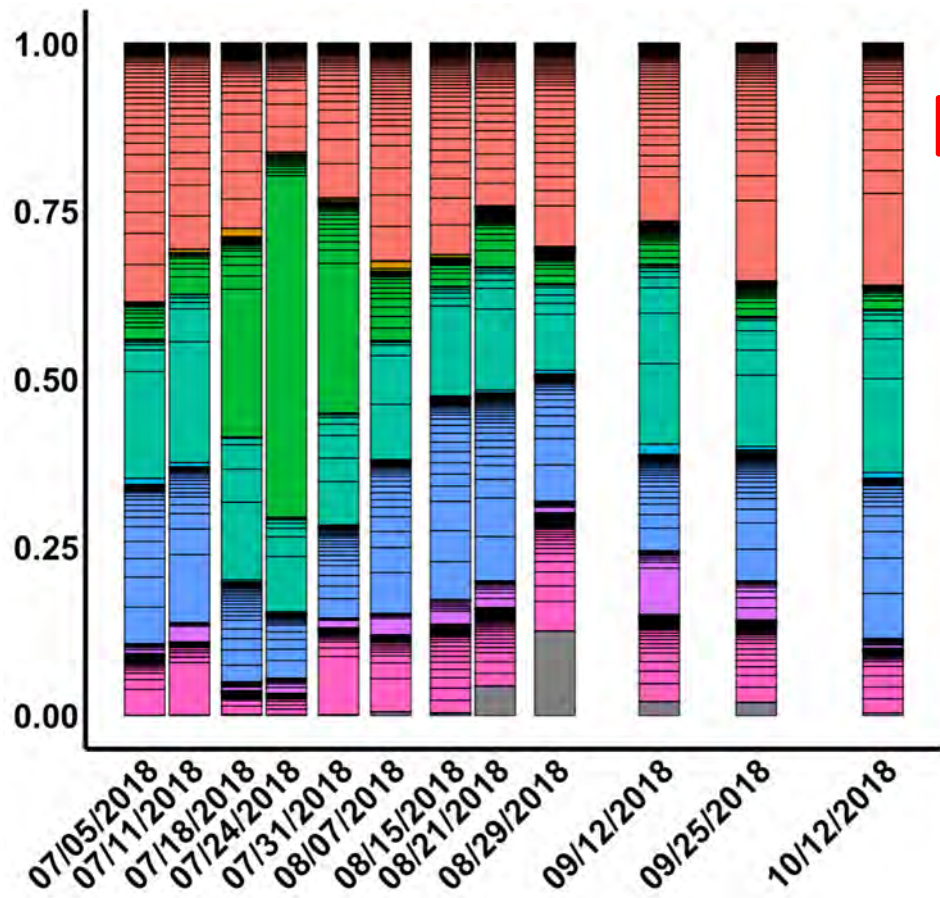
Next Steps



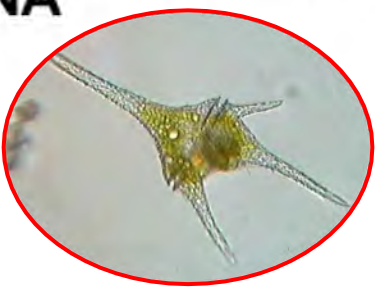
3 meter Secchi
Highland Lake
8/3/2022



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



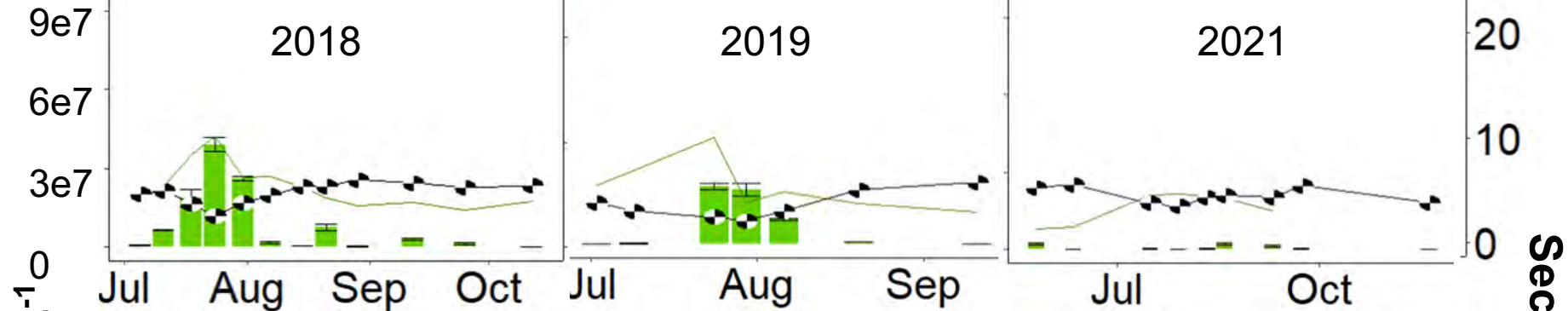
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- NA



Thank you,
Please send additional questions to:
sharon.mann@maine.edu
karen.wilson@maine.edu
rsleith@bigelow.org



Rhexinema (Green algae)



Cyanobium (Cyanobacteria)

