

ANZ Cyanobacteria Workshop 2023

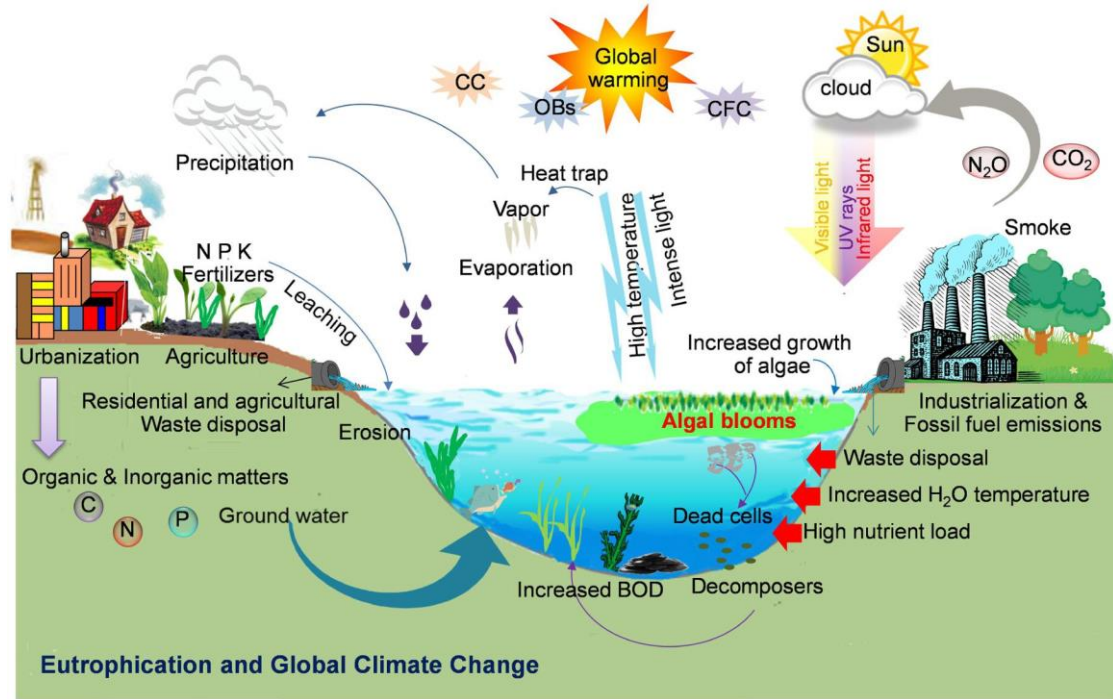
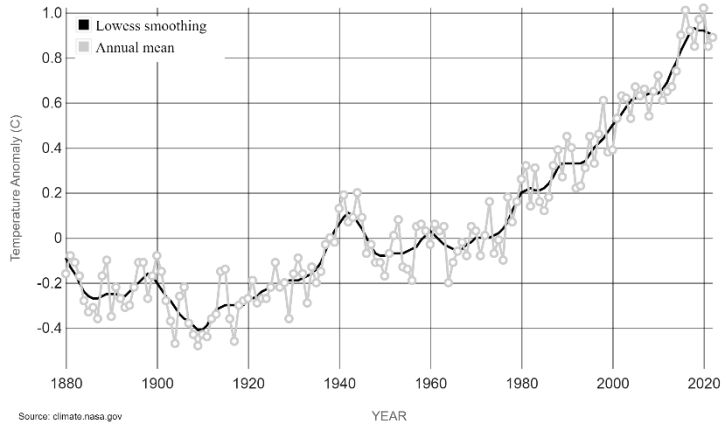
Metagenomic insights into Cyanobacterial blooms triggered by climate warming in NZ's freshwater lakes

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Response of freshwater systems to climate change

Trend in Global Temperature Rise



Reports on Cyanobacteria and Cyanotoxins in NZ Lakes

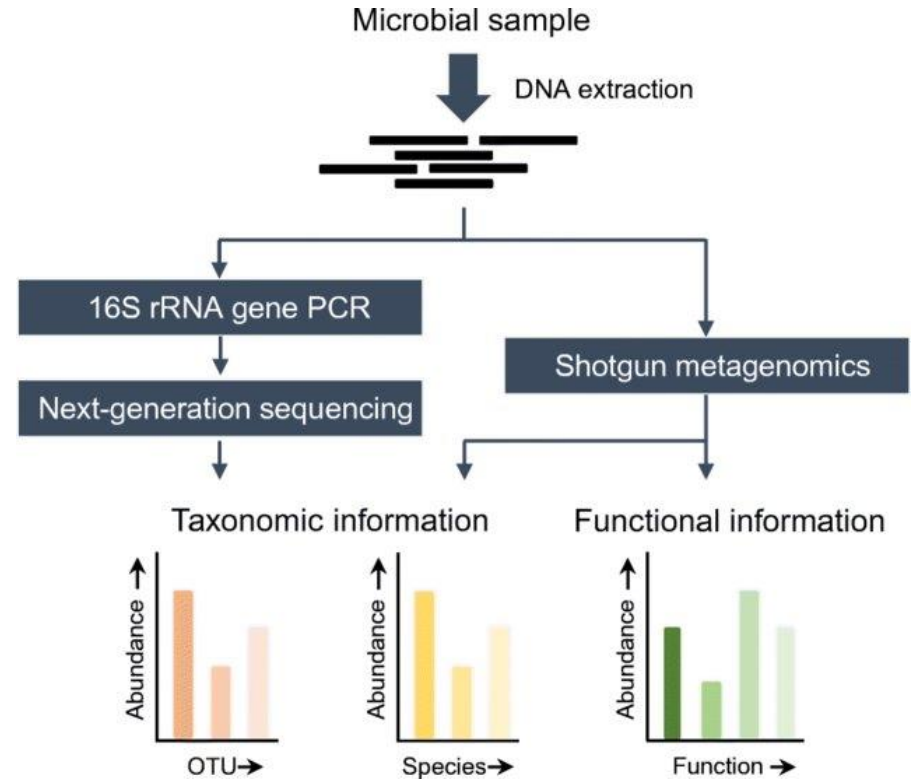
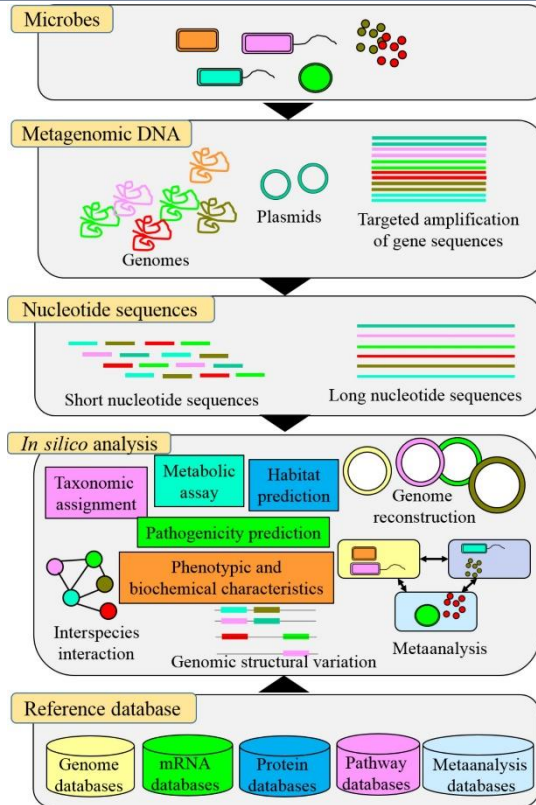
Table 1: Cyanobacterial genera known to occur in New Zealand fresh waters and the toxins they are known to produce worldwide.^{5,9}

Genera	Cyanotoxins known to be produced
<i>Anabaena</i>	anatoxin-a, ^a anatoxin-a(S), microcystins, ^a saxitoxins, ^a cylindrospermopsin
<i>Anabaenopsis</i>	microcystins
<i>Aphanocapsa</i>	microcystins
<i>Aphanizomenon</i>	anatoxin-a , cylindrospermopsin, saxitoxins, microcystins
<i>Arthrospira</i>	microcystins, anatoxin-a
<i>Cylindrospermum</i> ^b	cylindrospermopsin
<i>Cylindrospermopsis</i>	cylindrospermopsin , saxitoxins
<i>Lyngbya</i>	aplysiatoxins, lyngbyatoxin-a , saxitoxins
<i>Microcystis</i>	anatoxin-a, cylindrospermopsin, microcystins , saxitoxins
<i>Nodularia</i>	nodularin
<i>Nostoc</i>	microcystins ^a
<i>Oscillatoria</i>	anatoxin-a , aplysiatoxins, microcystins, ^a anatoxin-a(S), homoanatoxin-a
<i>Phormidium</i>	microcystin, ^a anatoxin-a , homoanatoxin-a
<i>Planktothrix</i>	microcystins, ^a homoanatoxin-a, anatoxin-a, aplysiatoxins, saxitoxins
<i>Pseudanabaena</i>	microcystins
<i>Raphidiopsis</i>	cylindrospermopsin, anatoxin-a, homoanatoxin-a, microcystins
<i>Snowella</i>	microcystins
<i>Synechocystis</i>	microcystins
<i>Woronichinia</i>	microcystins

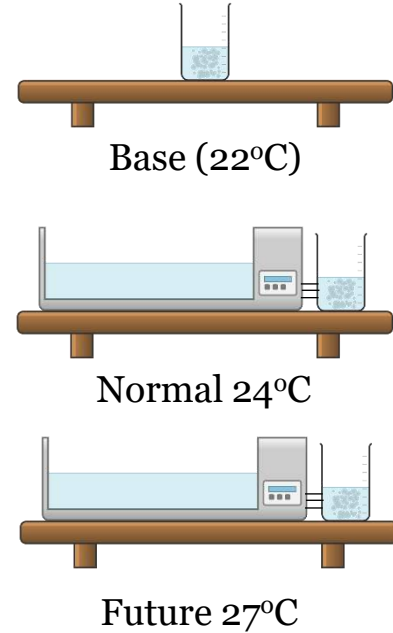
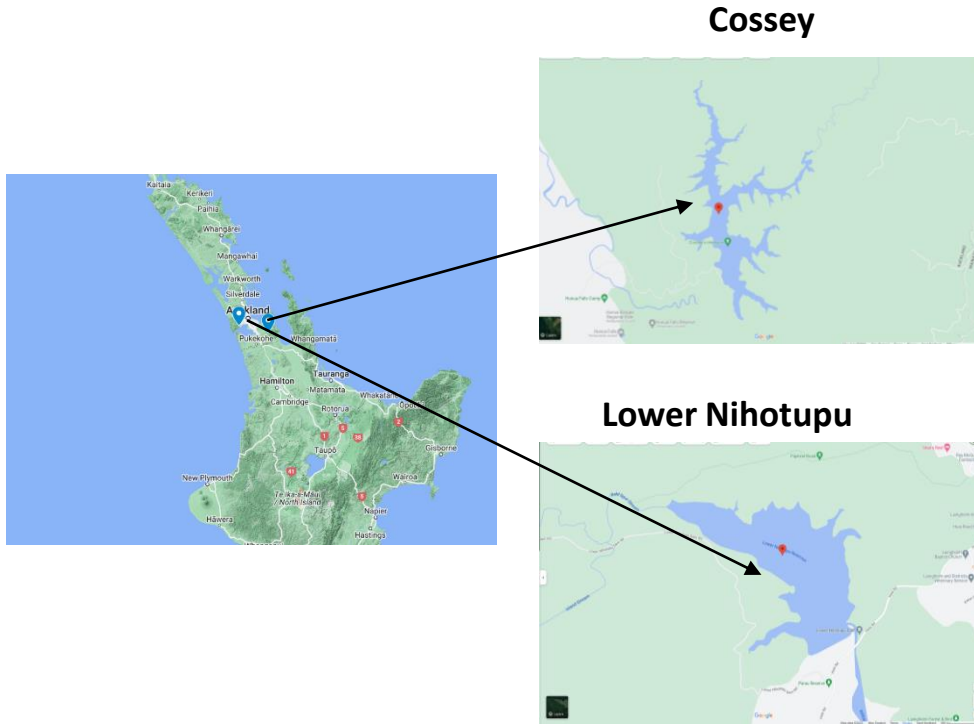
Current Cyanotoxin detection approach:

- Key cyanobacterial gene (*e.g.*, *mcyE*, *anaC*, *cyrJ*, *sxtA* etc.) detection through qPCR techniques.
- Detection of cyanotoxins (*e.g.*, Anatoxins, dihydro-anatoxins, Cylindrospermopsin, Microcystins/Nodularin, and Saxitoxins etc.) using LC-MS.

Metagenomics: Who are they? What are they doing?

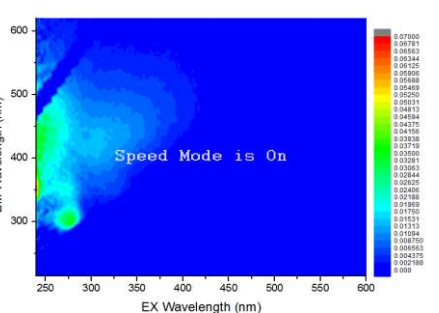
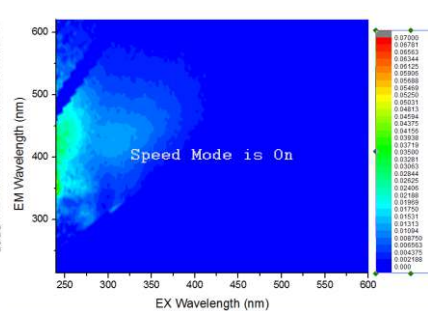
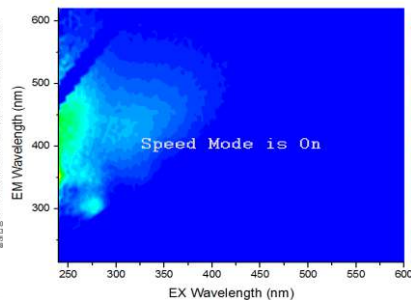
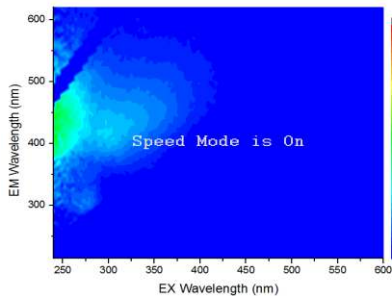


Simulating Climate warming induced microbial shifts

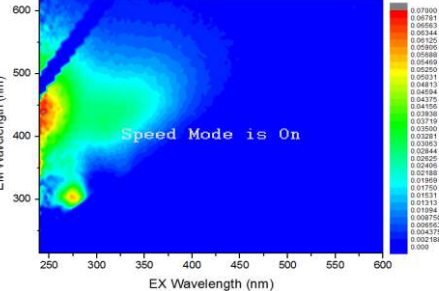
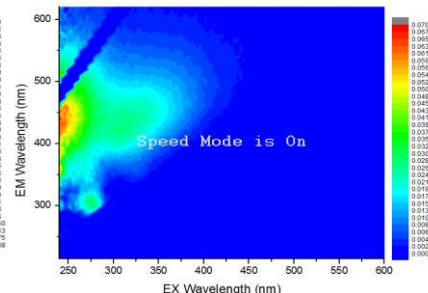
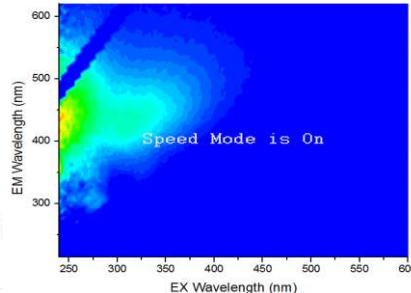
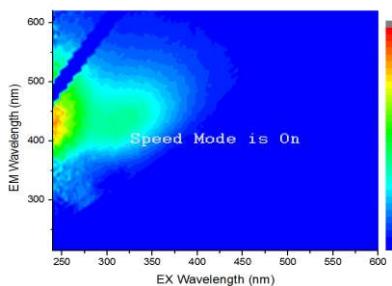


Variation in dissolved organic matter

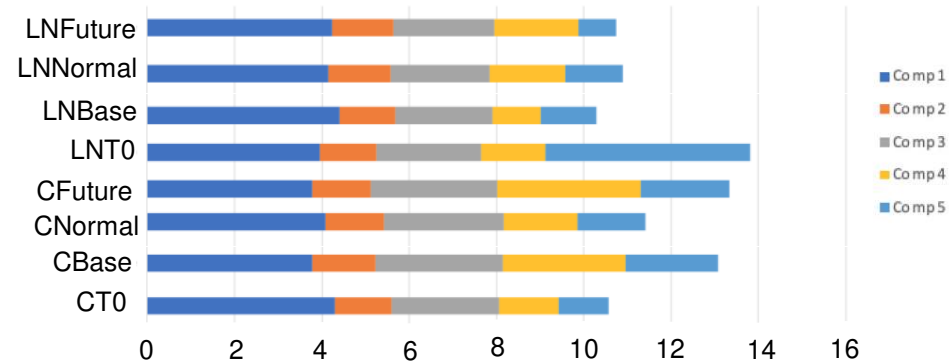
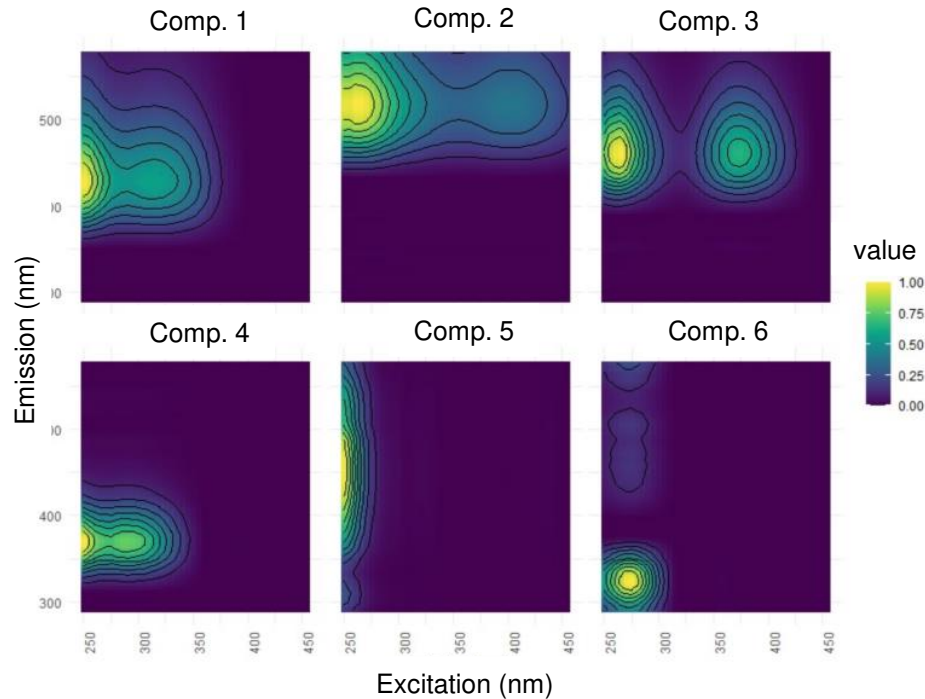
Cossey



Lower
Nihotupu

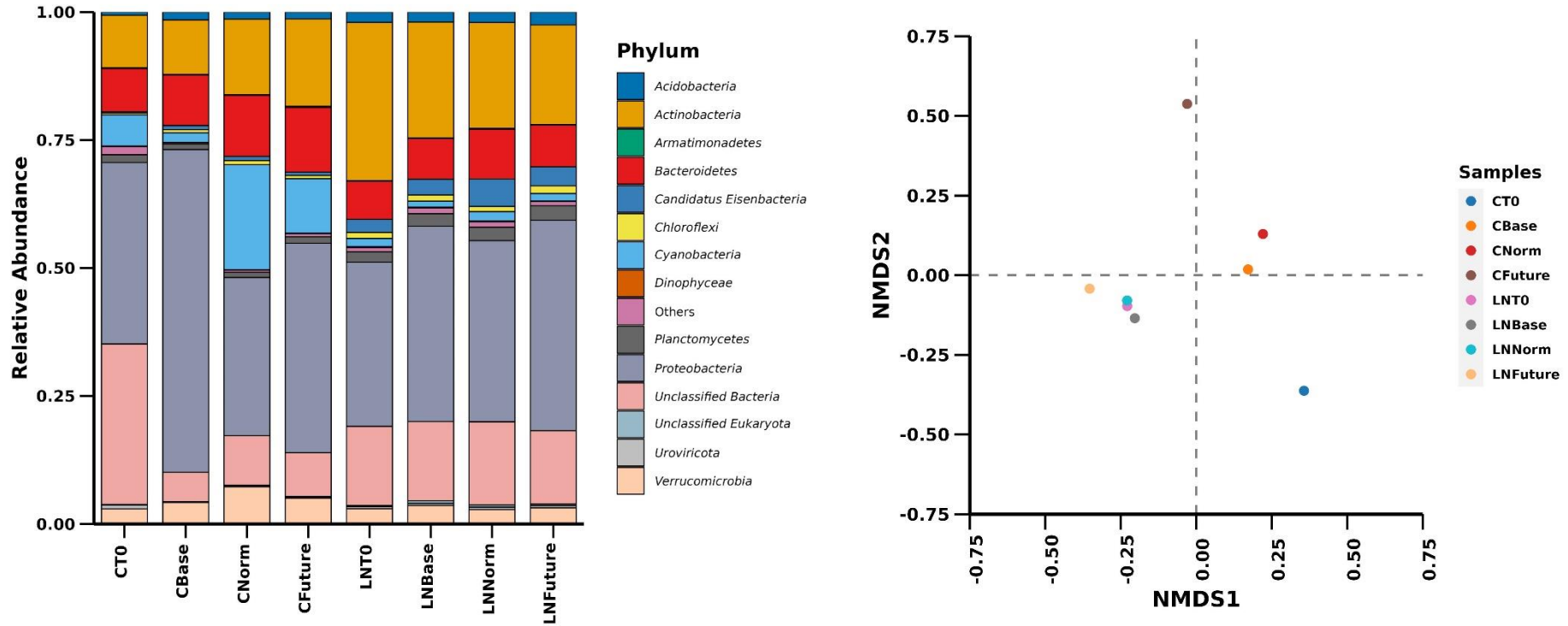


Variation in dissolved organic matter components

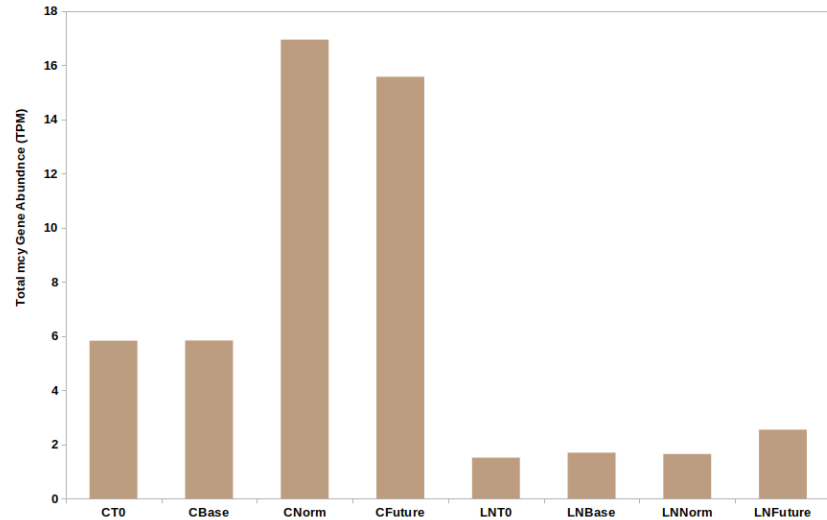
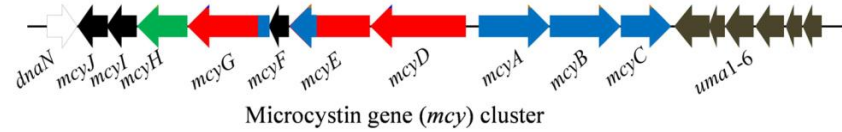
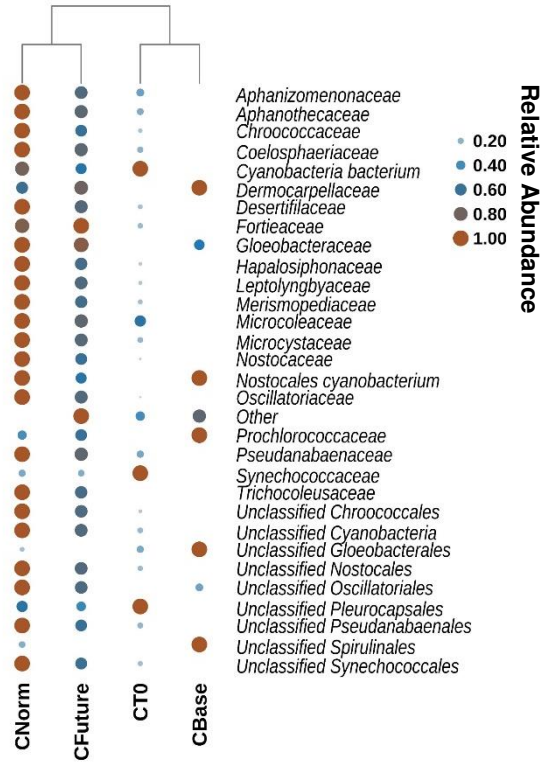


Component	Assignment
Comp. 1	Humic like Terrestrial
Comp. 2	Humic like Microbial derived
Comp. 3	Humic like
Comp. 4	Protein like Or Tryptophan-like

Climate warming induced shift in microbial communities



Cyanotoxin producers and total *mcy* gene Abundance



Summary

- ❑ Climate change poses significant risks to drinking water sources by promoting algal growth, pathogen proliferation, and toxin formation.
- ❑ Metagenomics analysis detects notable variations in microbial communities. Rising temperatures impact the population of cyanobacteria in Cosseys Reservoir.
- ❑ Warmer conditions lead to an increase in Cyanobacteria like *Microcystis*, *Nostoc*, *Anabaena*. The *mcy* gene clusters, linked to microcystin production, show an upward trend with higher temperatures.
- ❑ Elevated temperatures stimulate cyanobacterial pathways involving nitrogen, photosynthesis, and carbon fixation. This promotes toxin-producing organisms and amplifies toxin production.
- ❑ Findings shed light on climate change's potential influence on freshwater lakes with greater susceptibility to cyanobacterial blooms.
- ❑ Metagenomics based approach can enable early detection of harmful blooms in freshwater systems.

Acknowledgements



Emma Jay

NS Group



Funding Body

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High-performance Computing and Storage Facility



International Relations



Any
Question

