

Investigation into the presence of cyanobacteria in the **Quinnipiac River**

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Introduction

The area surrounding the Quinnipiac River is known to have an assortment of point and nonpoint pollution sources that have led to increased levels of nutrients and bacterial pollution (QRWA 2013). Elevated levels of nutrients can result in an excess growth of a single type of algae, known as an algal bloom. Urban development can cause a rise in nutrients, and therefore can impact the frequency and extent of these blooms (Huisman et al. 2018). Blooms of toxin producing cyanobacteria species can cause particularly serious health complications in aquatic and land organisms, including humans (Codd et al. 1999). Toxic cyanobacteria have been identified throughout the state of Connecticut (CT DEEP 2011); however, little research has been done on the Quinnipiac River to determine the presence and potential toxicity of cyanobacterial species. This study aims to identify the presence of cyanobacteria in the Quinnipiac River, as well as nutrients and water qualities that might correlate with the distribution of these species. Through collection and isolation of individual cyanobacterial species we intend to identify cyanobacterial communities and their potential for toxicity within the river. Furthermore, information gathered in this study will improve understanding of bacterial and pollutant levels throughout the Quinnipiac River, and provide useful knowledge to improve water quality.

Results

- Spirogyra sp. 9 AS-2014 voucher ARS06711_00001 EF426579 , Spirogyra sp. 00509 23S ribosomal RNA gene M676813 . Spirogyra sp. 1 AS-2014 (M676752, Zygnema sp. 1 AS-2014 geotia sp. 1 AS-2014 voucher ARS06197 00001 23S Mougeotia sp. 2 AS-2014 voucher ARS06695_00001 23S DQ229107, Chara vulgaris chloroplast DQ422812, Chlorokybus atmophyticus chloroplast Chlorella vulgaris strain TMCC3 23S ribosomal RNA Chlorella vulgaris strain ESP-31 23S ribosomal RN Chlorella vulgaris strain TMCC6 23S ribosomal RN/ Chlorella vulgaris isolate CAU 23S ribosomal RNA hlorogonium elongatum strain TMCC11 23S ribosoma Chlorococcum tatrense culture UTEX:2227 chloroplas KT625090, Chlorogonium capillatum culture-collection UTEX



Figure 3. Image of AC1349-05, identified as Chlorella vulgaris, at 400x magnification.



Study Objectives

• Determine the presence and identification of cyanobacteria in the Quinnipiac River. • Identify patterns of nutrient distribution and potential correlations between nutrients and the presence of cyanobacteria.





Discussion

We have identified a broad range of algal species present in the Quinnipiac River, but no cyanobacterial species have been isolated in cultures to date. This could be due to the latency of cyanobacterial development, requiring additional time to accommodate culturing and growth. Alternatively, the absence of cyanobacteria isolates could be the result of significantly low levels present in the river. Confirmation of cyanobacteria in similar waterways throughout the state of Connecticut (CT DEEP 2011) indicates that additional research, with an emphasis on identifying potential toxin producing species, is needed. Further investigation of the samples collected during the May-August collection period is necessary to identify all species, including cyanobacteria, present. Examination of these samples will continue as mixed and single species cultures mature to provide a thorough community assessment. Additionally, water quality and nutrient patterns will be compared to fluctuations in the type and amount of species present in the river to determine any correlations. Current and future results will provide an understanding of the environmental community as a whole, as well as contribute to a broad range of reference sequences to support an overarching study of environmental sequences of the Quinnipiac River.

Figure 1. Benthic samples were collected by utilizing a toothbrush to completely clear a 4cm² plot into pure distilled water.

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References

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