



Abstract

Emerald ash borers (EAB) are an invasive species in the United States that specifically **target ash trees**. **Lichens**, plant-fungus symbiotic organisms, are often used as **health indicators for the trees they grow** on. We hypothesized that a specific species of lichen growth may correlate with EAB infestation. **DNA sequencing** of lichen and bark samples from infested ash trees was done to determine present lichen species. One sample was identified as Gallowayella weberi (Xanthomendoza weberi), a common type of lichen ubiquitous across much of the contiguous United States. Since trees in declining health tend to be suitable lichen substrates, this could suggest that lichen are useful indicators on ash trees for possible locations of future infestations.

Introduction

- Emerald Ash Borer (EAB) are an invasive species that target ash trees
 - Bore through bark and lay eggs which consume phloem and cambium of tree
- Identifiable by D-shaped exit holes
- EAB infestation has killed tens of millions of ash trees throughout the US and Canada
- Older-growth ash trees in Garden City are being cut down to curb the spread of infestation
- Lichen = mycobiont (fungi) + photobiont (algae or cyanobacteria)
- Does not negatively impact substrate
- Often used as marker of thriving environment
- Previous studies have suggested relationship between lichen growth and diseased trees
- \circ Decreased foliage \rightarrow increased sunlight \rightarrow increased lichen growth
- We hypothesize there may be a potential correlation between infested trees and the type of lichen growth







Figure 3 (left): Phylogenetic tree shows low genetic relations between samples.

Determining Correlation between Invasive Species Infested Trees and Lichen Growth in Garden City

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Materials and Methods

Sample Collection

- Collection location: Terrace Park (40.73°06'67", -73°67'00")
- Utilized direct hand collection method
- Tweezers, gloves, plastic bag
- 4-6 cm in length
- Sampled 3 EAB infested ash trees marked to be cut down
- Collected late March
- Stored in freezer until isolation

Bioinformatics

- DNA Sequences manually trimmed
- Samples identified on NCBI BLAST
- MUSCLE and Phylogenetic diagram constructed on DNA Subway



QSJ-002

QSJ-004/QSJ-005

QSJ-006

Figure 1 (above): EAB infested ash trees on Terrace Park where samples were collected 3/28/23. Trees are marked to be cut down due to infestation.

Figure 4 (right): Comparison of samples using MUSCLE diagram shows poor quality sequencing (frequent gray areas) except for QSJ-005.



Isolating the DNA

- Utilized Silica DNA extraction method
- Using tweezers, 1 cm pieces of samples were deposited into Eppendorf tubes
- Lysis solution and pestle used to mechanically break membranes and release DNA
- Supernatant was separated into another tube and silica resin was pipetted in
- Silica resin's polarity allowed it to bind to DNA • Silica pellet was washed with wash buffer 2 times
- dH₂O mixed with pellet to dissolve
- Fungi (lichen specific) ITS-1 primer amplified using CSHL protocol
- Gel electrophoresis performed to confirm success of
- Samples sent to Cyverse to obtain DNA sequences

ith SYBR → QSJ-0 ∞ QSJ-00 ဖ QSJ-006

Figure 2 (above): PCR products were identified using Gel Electrophoresis. Lanes 4-9 contain samples; odd wells contain tree bark, even wells contain lichen. QSJ-002, QSJ-004, QSJ-005, and QSJ-006 show positive results. QSJ-002, QSJ-004, and QSJ-006 suggest double bands (see discussion).

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• QSJ-005 (bark) likely yielded lichen due to trace bits of lichen left on bark amplified by ITS fungi primer; succeeded over intended lichen samples because less fungi present • Future Directions • Collect lichens from non-infested ash trees to



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Discussion

• Sample QSJ-005 was identified as *Gallowayella* weberi, suggesting there may be a correlation between EAB infestation and Gallowayella weberi • 3 out of 4 sequenced samples had too low quality to be identified on BLAST

- QSJ-002, QSJ-004, and QSJ-006 in gel show double bands \rightarrow suggests multiple fungi \circ Multiple species of fungi \rightarrow background noise in sequencing, no dominant fluorescence signal detected \rightarrow low-quality sequence/ unable to identify
 - compare lichen species
- Correlate severity of infestation with lichen amount
- Decreased foliage in infested trees may lead to increased lichen growth
- \circ Bark DNA samples still remain \rightarrow use insect specific primer to confirm EAB presence

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