

# Environmental DNA surveillance for aquatic plants: a 21<sup>st</sup> century tool for species detection, biodiversity monitoring and management

Chris Wilson and Stephanie Coghlan

Aquatic Research and Monitoring Section,

Ontario Ministry of Natural Resources and Forestry



# Environmental DNA (eDNA)

cells / DNA shed from living or dead organisms into surrounding environment

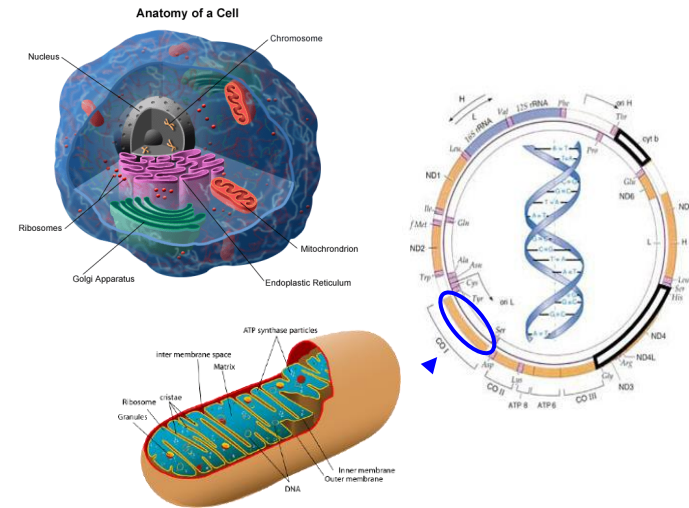
“molecular smoke alarm”

doesn't say what the source was

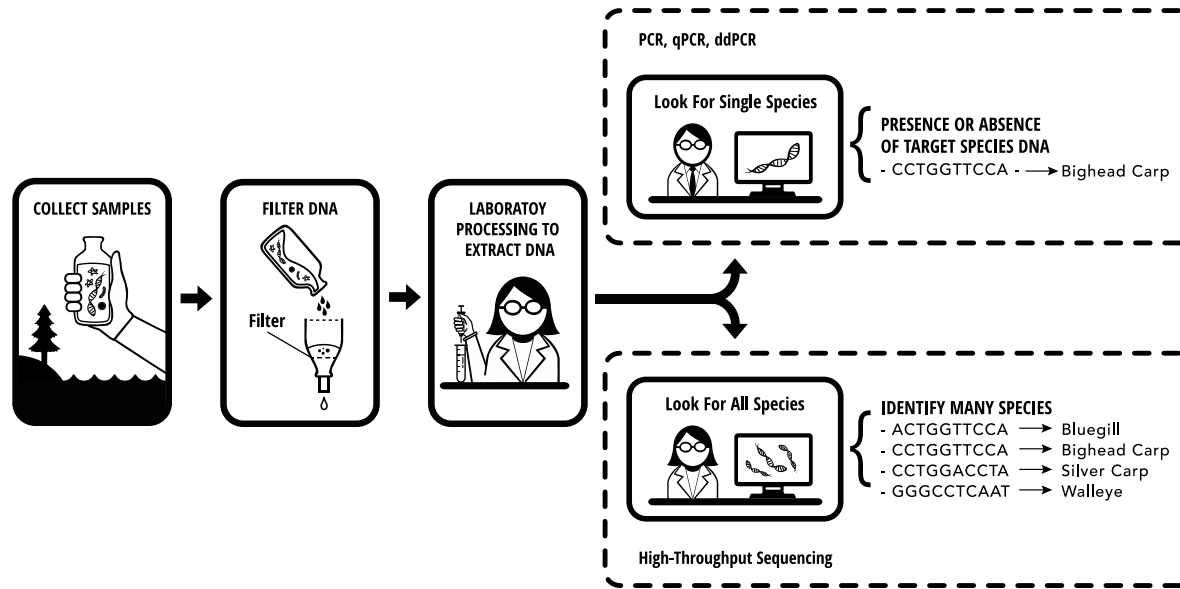
- living / dead; body parts; fluids, cells, free molecules, etc.

doesn't tell re: age, sex, size, reproductive status, population info

➤ **positive detection** = DNA from the species was present at that location when the sample was collected



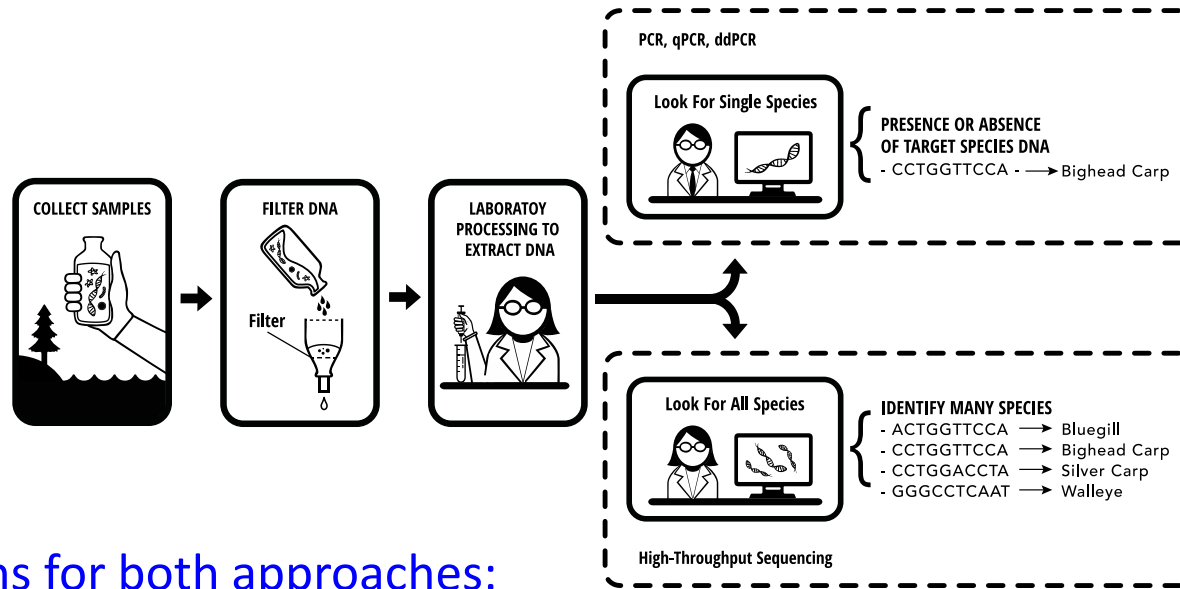
# Complementary approaches: targeted and community eDNA assays



Active surveillance: **targeted single species** eDNA testing to evaluate if a species is present (bighead carp, tench)

Passive surveillance: use **community eDNA** or **eDNA metabarcoding** to characterize aquatic communities / local species assemblages (fish, invertebrates, plants, etc.) using high-throughput sequencing

# Complementary approaches: targeted and community eDNA assays

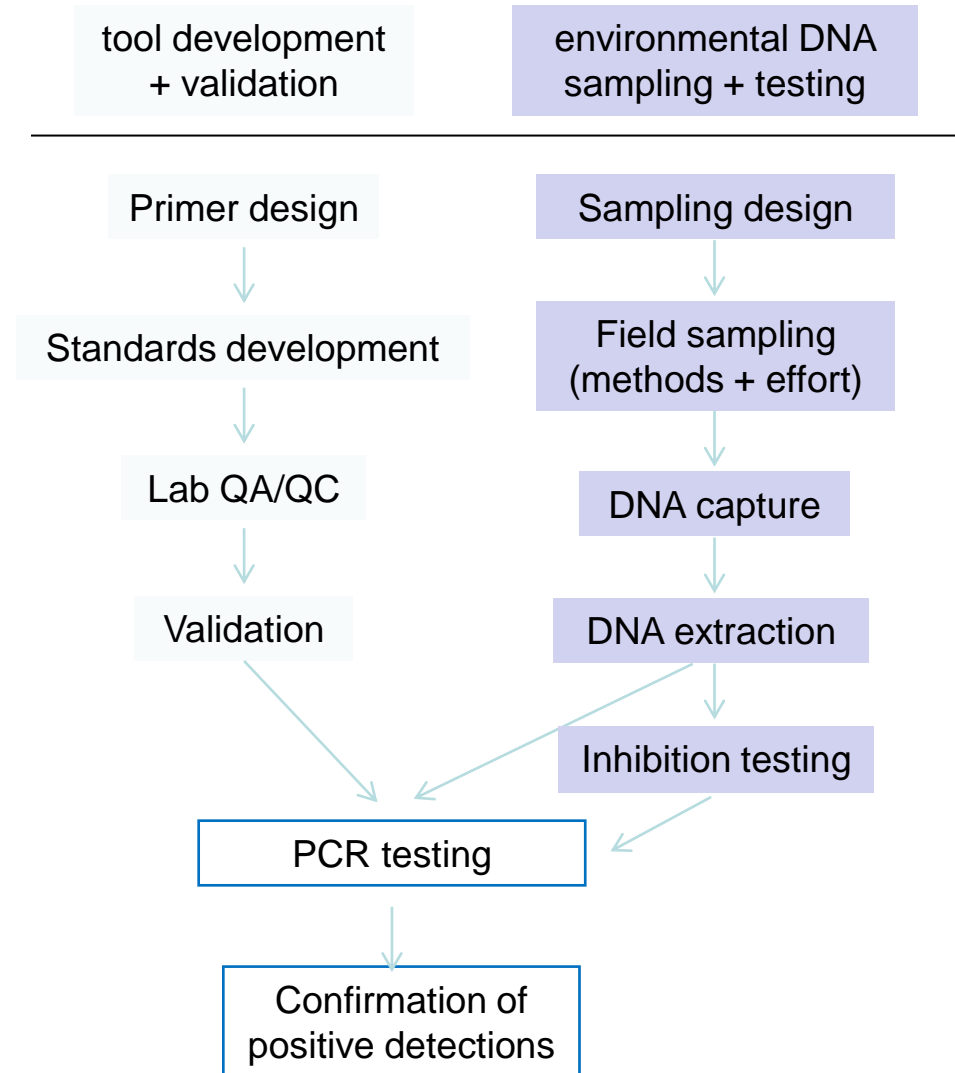


## Considerations for both approaches:

- species specificity and sensitivity
- spatial sensitivity
- temporal sensitivity
- quantitative sensitivity

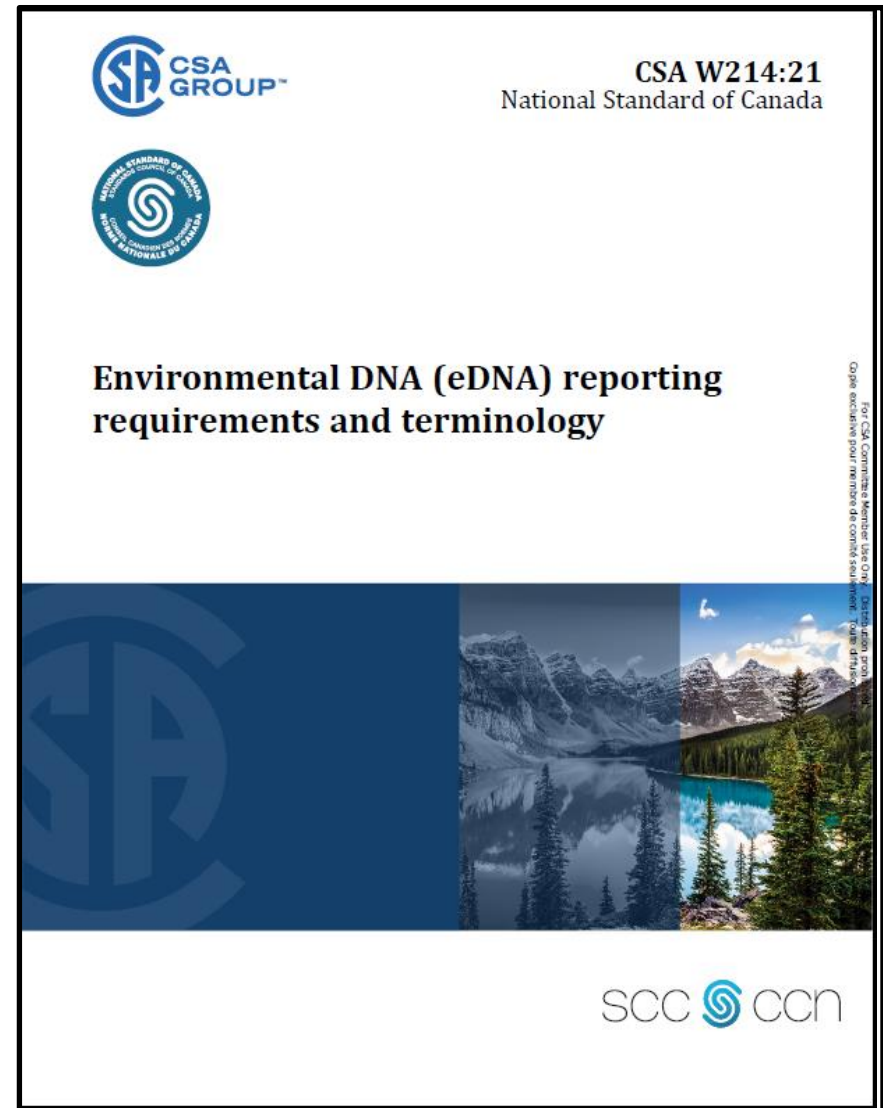
# Information needs for eDNA data acceptance

- “Invisible Man” scenario
- habitat description + characteristics
- sampling design + effort
- spatial + temporal metadata
- assay metadata
  - technology used
  - test conditions (details)
  - assay sensitivity + specificity
  - replicates + controls



# Information needs for eDNA data acceptance

- “Invisible Man” scenario
- habitat description + characteristics
- sampling design + effort
- spatial + temporal metadata
- assay metadata
  - technology used
  - test conditions (details)
  - assay sensitivity + specificity
  - replicates + controls



([www.csagroup.org](http://www.csagroup.org))

# Challenges for aquatic plant eDNA

- more challenging DNA barcoding for assay design
- less available baseline sequence data
- lower shedding rate vs animals (movement, excretion, temperature-related activity levels, vegetative reproduction)
- seasonal variation in shedding (growth vs. senescence)
- long-distance drifting of plant remains (rafting, etc.)
- passive dispersal by boats, trailers, propellers
- dormant life stages



# Targeted eDNA assays for “least wanted” aquatic plants

Aquatic Botany 122 (2015) 27–31



Contents lists available at ScienceDirect

Aquatic Botany

journal homepage: [www.elsevier.com/locate/aquabot](http://www.elsevier.com/locate/aquabot)



Short communication

Development of species-specific environmental DNA (eDNA) markers for invasive aquatic plants



Michelle Scriver<sup>a</sup>, Allison Marinich<sup>b</sup>, Chris Wilson<sup>c</sup>, Joanna Freeland<sup>a,\*</sup>

<sup>a</sup> Department of Biology, Trent University, Peterborough, ON K9K 1X4, Canada

<sup>b</sup> Environmental and Life Sciences Graduate Program, Trent University, Peterborough, ON K9K 1X4, Canada

<sup>c</sup> Aquatic Biodiversity and Conservation Unit, Ontario Ministry of Natural Resources and Forestry, Trent University, Peterborough, ON K9K 1X4, Canada



matK, rbcL, trnH-*bsbA* as target sequences:

Water soldier

Eurasian water milfoil

Carolina Fanwort

Parrotfeather

Water hyacinth

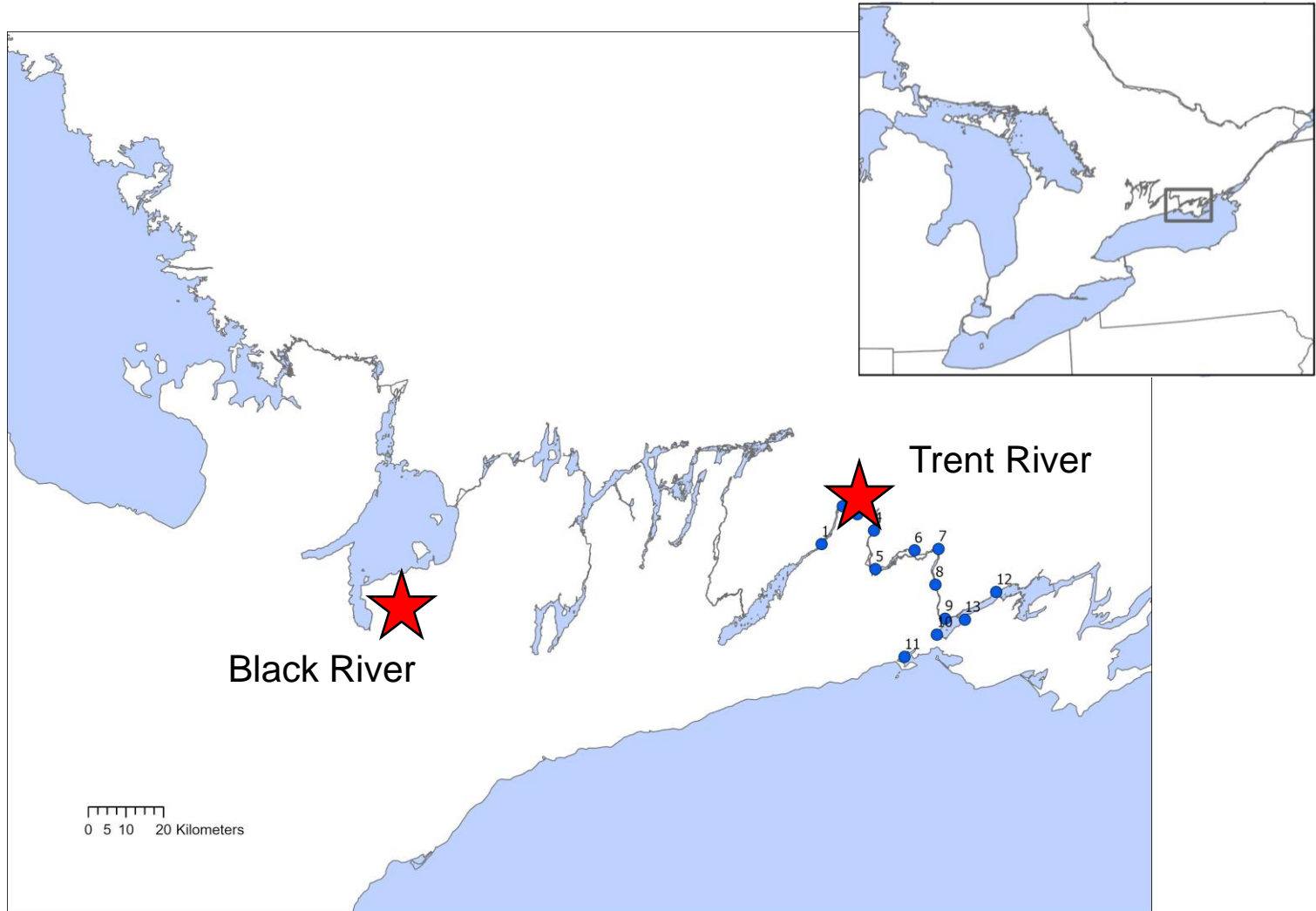
Water lettuce

Yellow floating heart

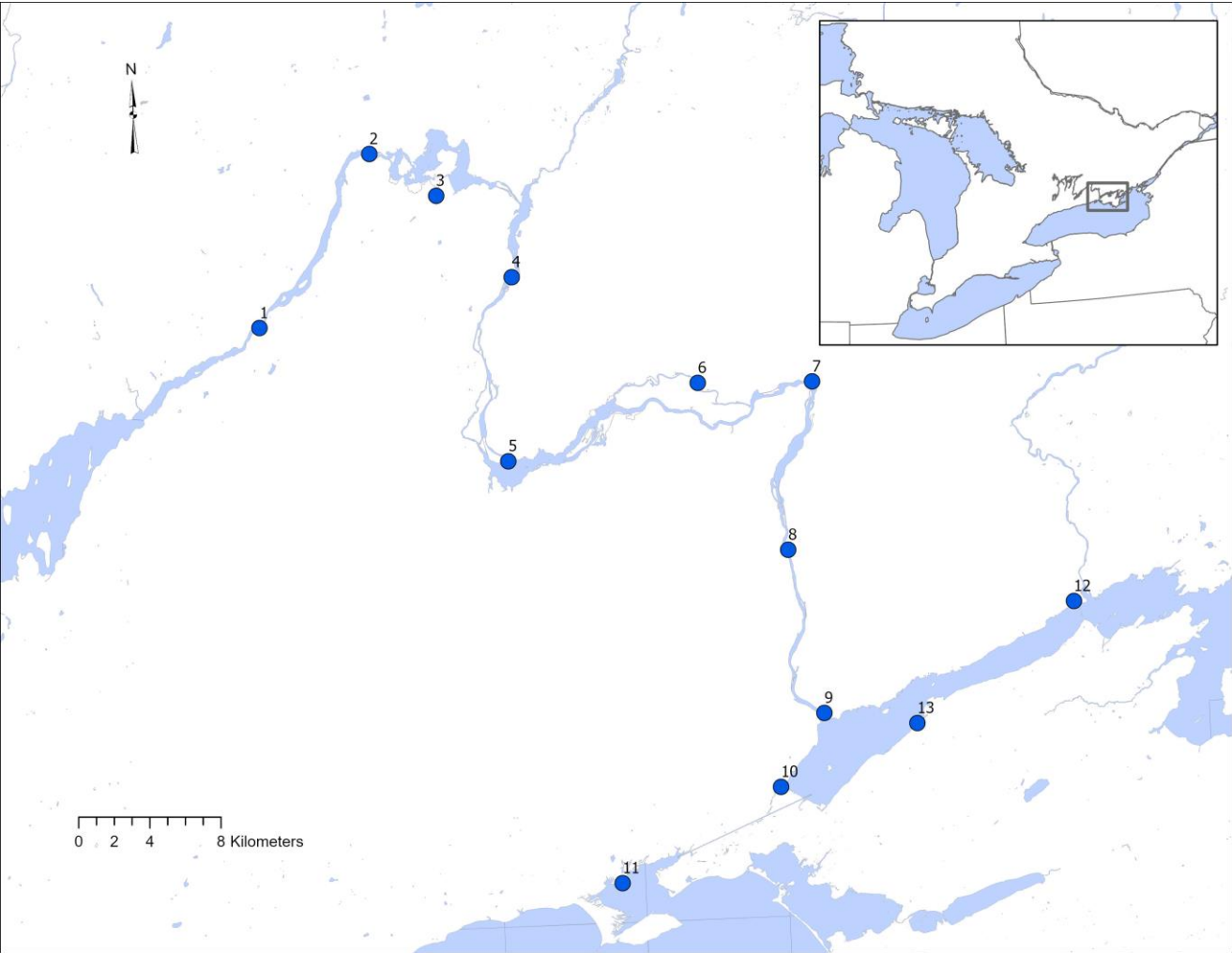
*(Scriver et al. 2015)*



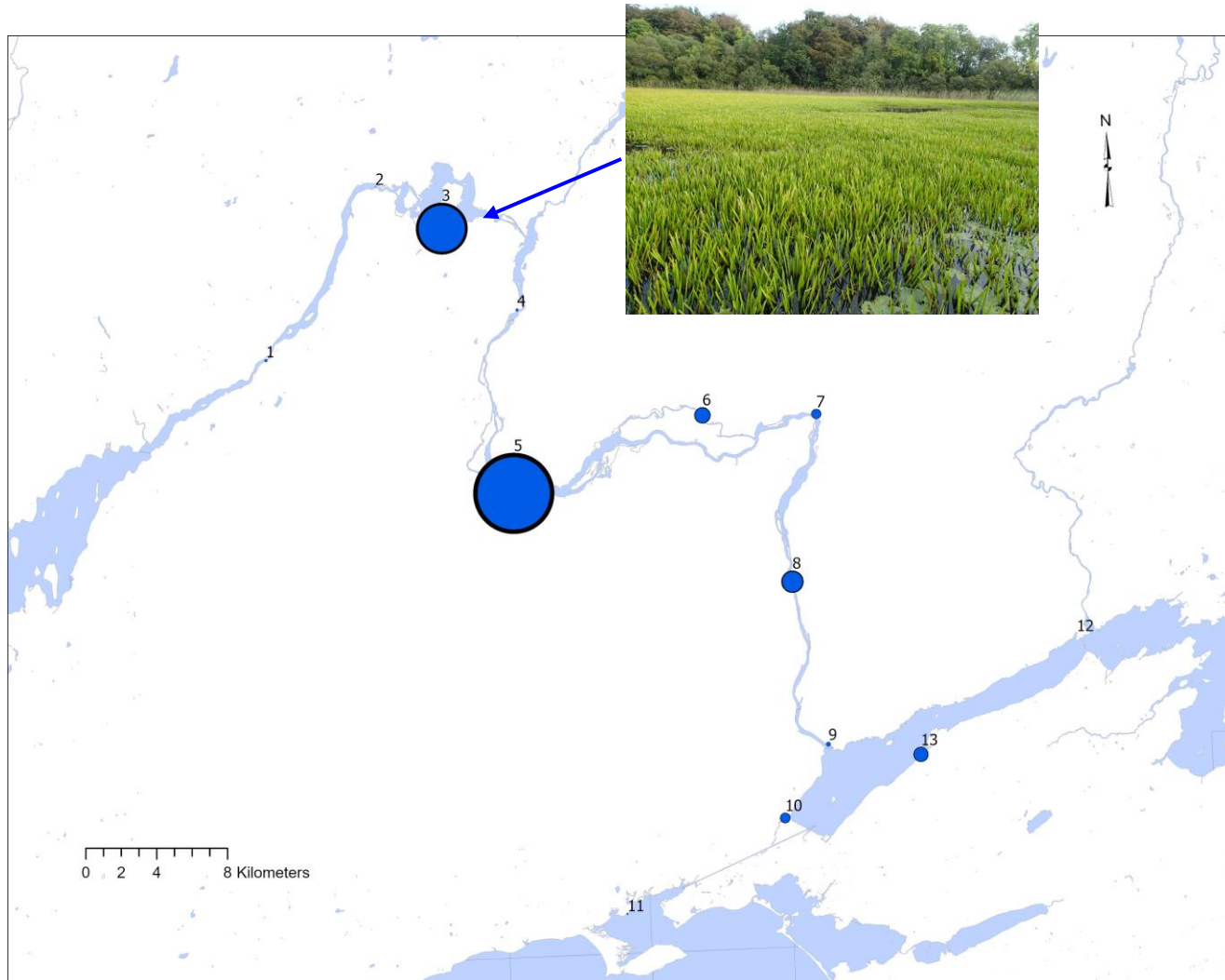
# Water soldier infestations in Ontario



# Water soldier eDNA sampling in the Trent-Severn Waterway

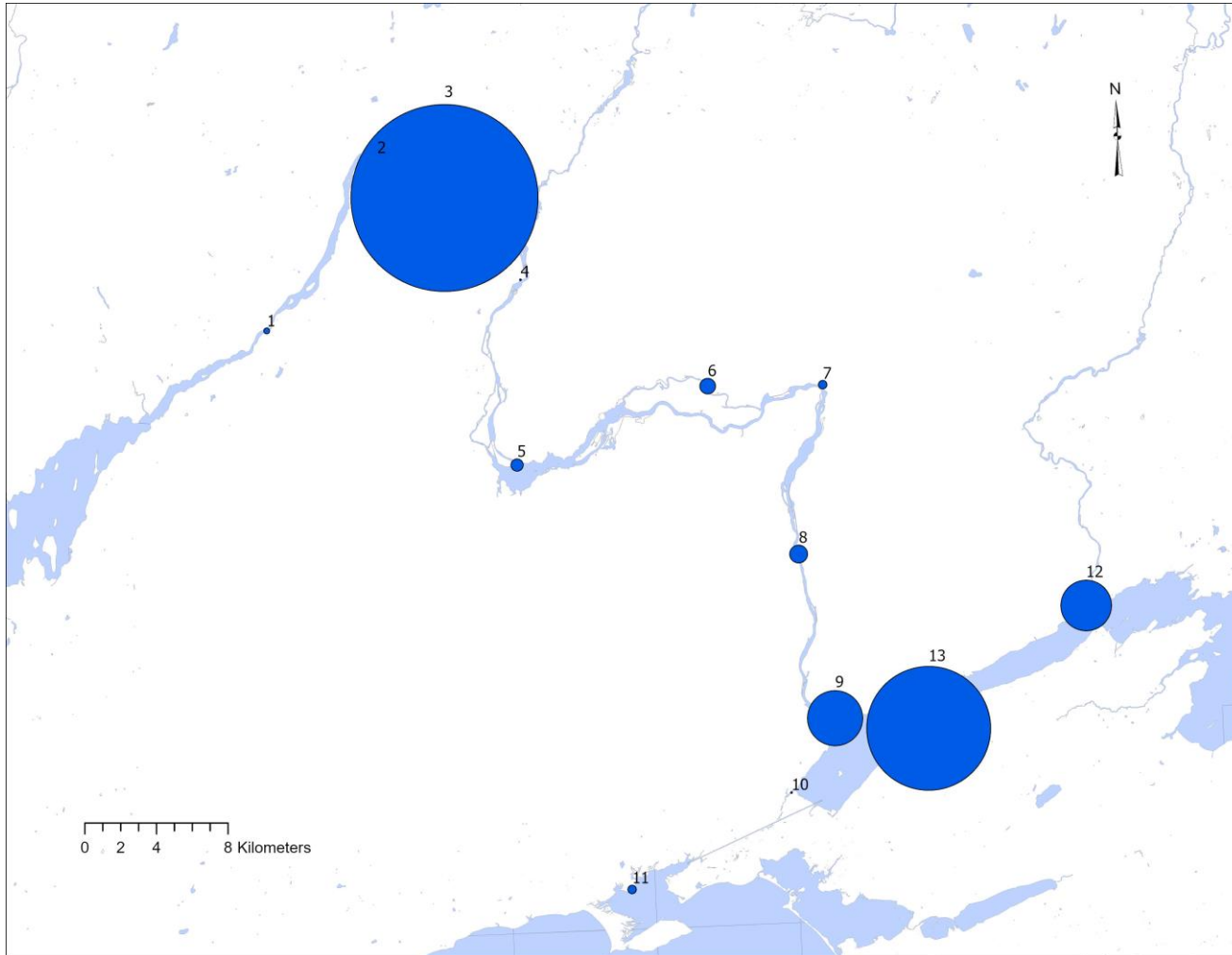


# September 2021 eDNA detection results (pre-herbicide treatment)



- strong detections at known established populations
  - weak detections at sampling sites without known species presence, including Bay of Quinte in Lake Ontario
- *presence confirmed in 2022*

# November 2021 eDNA detection results (post-herbicide treatment)

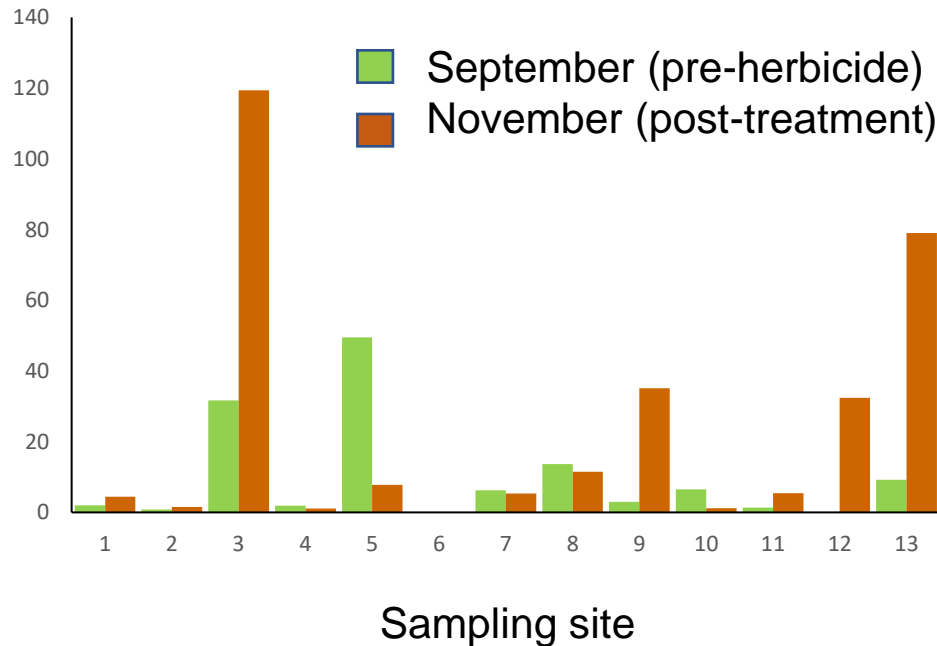


- very strong detection at treatment site
  - strong detections also at untreated sites in Bay of Quinte
  - weak detections at sites without known species presence at the time
- *presence confirmed in 2022*

# Herbicide treatment substantially affected eDNA results



*E. stratiotes* eDNA detection (copies/reaction)



## *Pre-herbicide treatment*

- weak detections upstream of established population (site 3) before herbicide treatment
- weak detections in Bay of Quinte and Presqu'île Bay (Lake Ontario)

## *Post-treatment*

- strong detections in treatment area
- Strong detections downstream of treatment area most likely reflect flushing of dead material

# Targeted eDNA efforts for invasive aquatic plants

## *Water soldier control / eradication*

- 2022 detections in Bay of Quinte visually confirmed by plant surveys
  - Local removal efforts
  - Expanded survey in 2023 incorporating hydrodynamic modelling
- Black River eradication considered successful after 3 years of no positive detections

## *Other species*

- Water chestnut (Kingston area, Lake Ontario) strong detections in treatment area
- Hydrilla (Niagara River)

## *Stakeholders / citizen science*

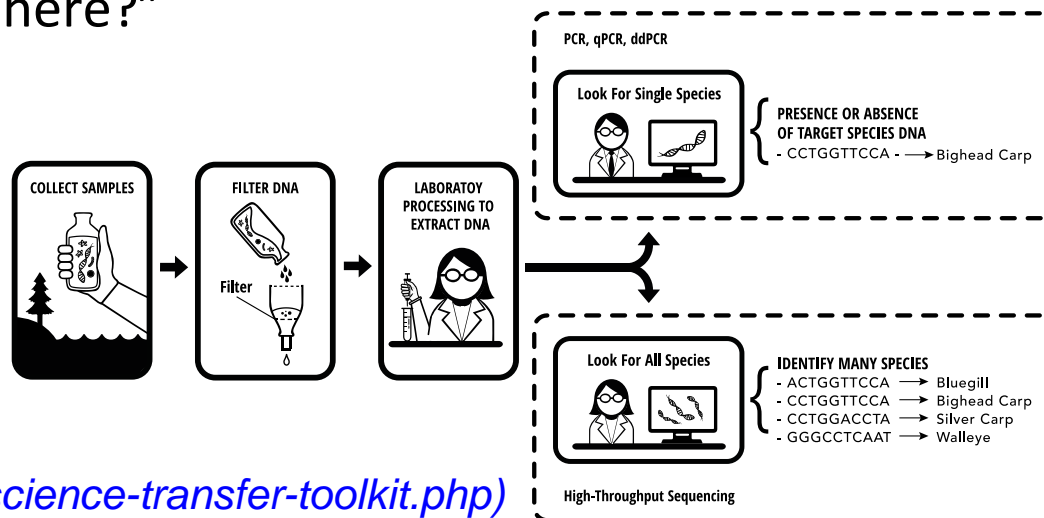
- Invasive Species Centre: eDNA sampling for community AIS detection in Ontario  
[www.invasivespeciescentre.ca/eDNA](http://www.invasivespeciescentre.ca/eDNA)



**Invasive Species Centre**

# Why eDNA metabarcoding?

- Large areas to cover – don't need to locate individual plants
- Experts required to ID physical specimens
- Species-specific eDNA solves some of these, but requires prior knowledge of expected species/species of interest
- Metabarcoding allows us to gain comprehensive knowledge of what species are present
  - Instead of “is water soldier here?” we're asking “what plants are here?”



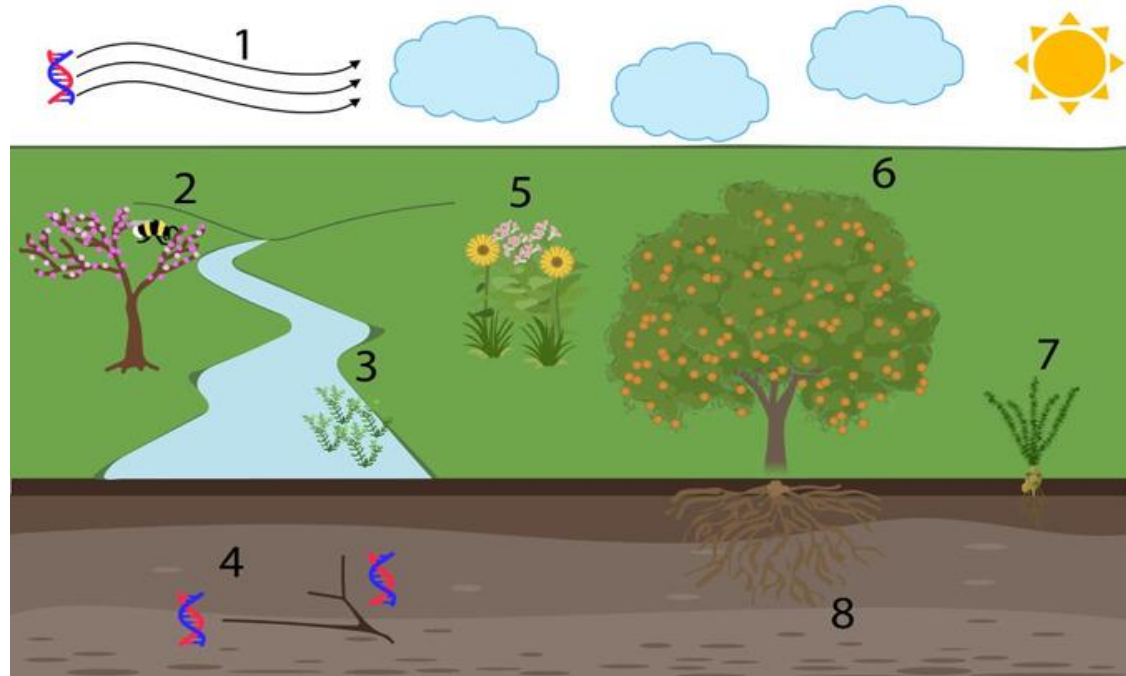
# Metabarcoding

Well-developed assays and databases for other organisms

- Animals, invertebrates, microbes

Sources:

- Water
- Soil
- Pollen
- Air



American J of Botany, Volume: 110, Issue: 2, First published: 11 January 2023, DOI: (10.1002/ajb2.16120)



# My MSc thesis

- Dr. Joanna Freeland & Dr. Aaron Shafer



- Develop aquatic plant metabarcoding assays
- Developed assays for 3 gene regions
  - Well-represented in database and appropriate for metabarcoding
  - matK, rbcL, and ITS2
- Tested against metabarcoding assays (rbcL and ITS2) developed for terrestrial plant metabarcoding from soil samples (Fahner et al. 2016)

# Mock community

- eDNA sample of known composition
- Provides baseline of what to expect in results

**omatK2:** 9 species-level, 11 genus-level

**orbcl2:** 7 species-level, 17 genus-level

**omatK2 + orbcl2 + oITS2:** 14 species-level, 10 genus-level

Species	omatK2	orbcl2	orbclA	oITSn	oITS2
<b>Nymphaeales</b>					
<i>Cabomba caroliniana</i>					
<i>Nuphar variegata</i>					
<i>Nymphaea odorata</i>					
<b>Poales</b>					
<i>Phragmites australis</i>					
<i>Typha latifolia</i>					
<i>Typha minima</i>					
<b>Alismatales</b>					
<i>Elodea canadensis</i>					
<i>Hydrocharis morsus-ranae</i>					
<i>Potamogeton crispus</i>					
<i>Potamogeton strictifolius</i>					
<i>Stratiotes aloides</i>					
<i>Vallisneria americana</i>					
<b>Other monocots</b>					
<i>Acorus calamus</i>					
<i>Eichhornia crassipes</i>					
<i>Iris pseudacorus</i>					
<i>Pistia stratiotes</i>					
<i>Pontederia cordata</i>					
<i>Schoenoplectus acutus</i>					
<b>Dicots</b>					
<i>Ceratophyllum demersum</i>					
<i>Megalodonta beckii</i>					
<i>Myriophyllum aquaticum</i>					
<i>Myriophyllum sibiricum</i>					
<i>Myriophyllum spicatum</i>					
<i>Nymphoides peltata</i>					
<i>Trapa natans</i>					

*Coglan et al. 2021*

# Mock community

## AIS/species of concern in Ontario:

Carolina fanwort

Frogbit

Water soldier

Water hyacinth

Yellow iris

Water lettuce

Parrot feather

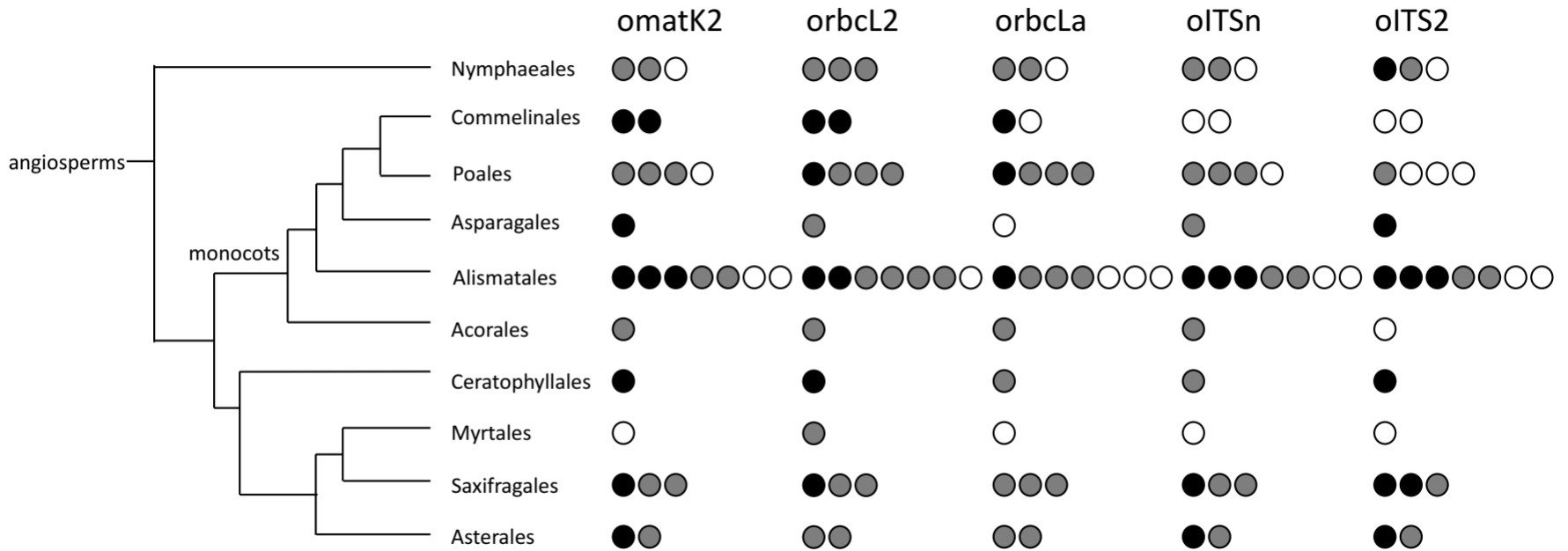
Eurasian watermilfoil

Yellow floating heart

European water chestnut

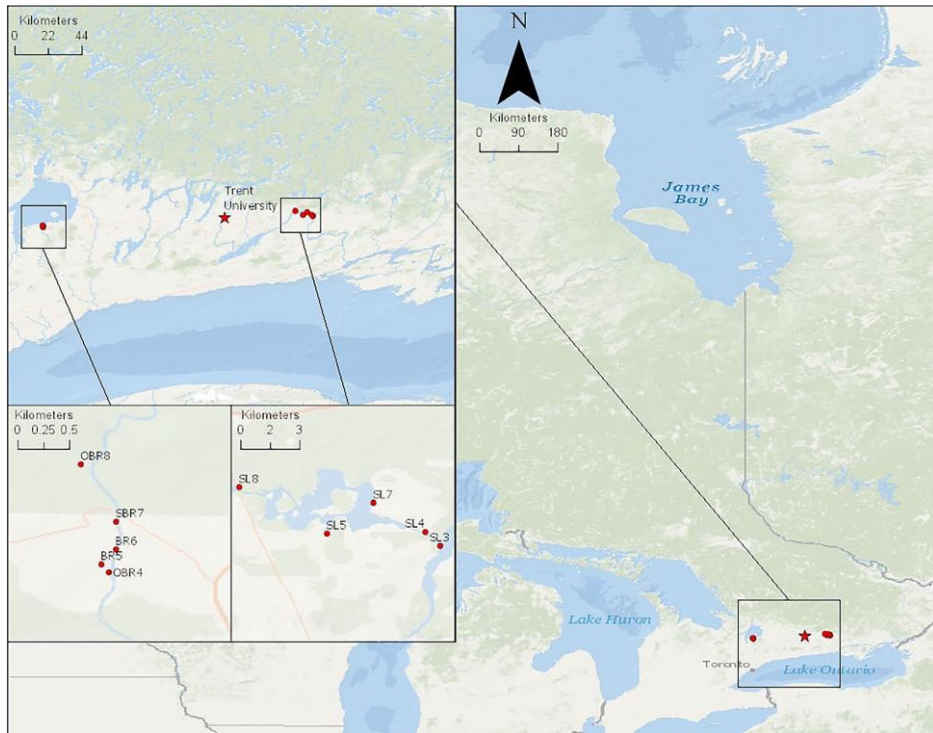
Species	omatK2	orbcl2	orbclA	oITSn	oITS2
<b>Nymphaeales</b>					
<i>Cabomba caroliniana</i>					
<i>Nuphar variegata</i>					
<i>Nymphaea odorata</i>					
<b>Poales</b>					
<i>Phragmites australis</i>					
<i>Typha latifolia</i>					
<i>Typha minima</i>					
<b>Alismatales</b>					
<i>Elodea canadensis</i>					
<i>Hydrocharis morsus-ranae</i>					
<i>Potamogeton crispus</i>					
<i>Potamogeton strictifolius</i>					
<i>Stratiotes aloides</i>					
<i>Vallisneria americana</i>					
<b>Other monocots</b>					
<i>Acorus calamus</i>					
<i>Eichhornia crassipes</i>					
<i>Iris pseudacorus</i>					
<i>Peltia stratifolies</i>					
<i>Pontederia cordata</i>					
<i>Schoenoplectus acutus</i>					
<b>Dicots</b>					
<i>Ceratophyllum demersum</i>					
<i>Megalodonta beckii</i>					
<i>Myriophyllum aquaticum</i>					
<i>Myriophyllum sibiricum</i>					
<i>Myriophyllum spicatum</i>					
<i>Nymphoides peltata</i>					
<i>Trapa natans</i>					

# Diversity of identified species



*Coghlan et al. 2021*

# Pilot sites



Coghlan et al. 2021

- Black River
- 17 species
- 24 genera
- Seymour Lake
- 14 species
- 27 genera

## Noteworthy detections:

Flowering rush (*B. umbellatus*)  
Frogbit (*H. morsus-ranae*)  
Yellow iris (*I. pseudacorus*)  
Parrot feather (*M. aquaticum*)  
Yellow floating heart (*N. peltata*)  
Phragmites (*P. australis*)  
Water lettuce (*P. stratiotes*)  
Water soldier (*S. aloidess*)

# Work since MSc

Testing orbcl2 on more sites and more replicates

- 44 sites
- orbcl2 identified all but one species from mock community to at least genus-level, and identified largest number of species from pilot sites

Monitor water soldier presence following removal/herbicide treatments

General results

- Noteworthy aquatics
  - AIS: European frogbit (*Hydrocharis morsus-ranae*), flowering rush (*Butomus umbellatus*), water soldier (*Stratiotes aloides*)

# Limitations

## Challenges with plant DNA in general

- Assay design: interspecific variation does not consistently exceed intraspecific variation; multi-assay approach might yield best results
  - Hybrids
- Taxonomic resolution – often genus-level ID
- Doesn't offer insight to abundance
- Relies on complete, error-free reference database

# Strengths and next steps

## Monitor biodiversity over time

- Incl. species occurrences before and after AIS invasions, patterns of co-occurring organisms
- Beyond plants – same samples can be used for any organisms
- Establish a baseline for routine monitoring

Early signs of invasions, particularly when you don't know what species you expect to find

May lead to targeted inquiries



# eDNA resources



[www.csagroup.org](http://www.csagroup.org)

GLFC Science Transfer eDNA

<http://www.glfc.org/science-transfer-toolkit.php>

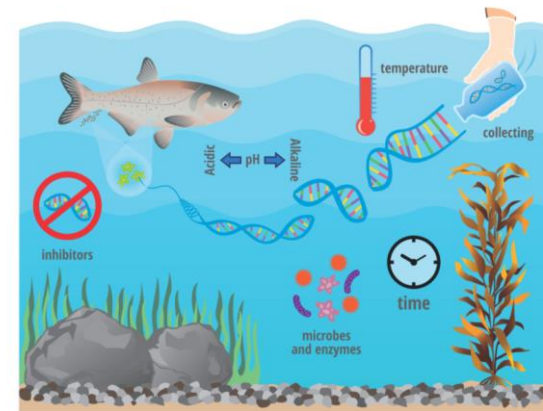


**GEN-FISH**

Genomic Network for Fish  
Identification, Stress and Health

Uses and Limitations of Environmental DNA (eDNA) in Fisheries  
Management

*Project leader: Welsh, A.*



[www.gen-fish.ca](http://www.gen-fish.ca)

eDNAtlas (USDA and USFS):

[www.fs.usda.gov/rmrs/projects/aquatic-ednatlas-project](http://www.fs.usda.gov/rmrs/projects/aquatic-ednatlas-project)