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**Environmental DNA (eDNA) Detection
of Aquatic Invasive Species in
Manitoba Lakes, a comparative analysis
of microscopy and
DNA detection methods.**

2026 Invasive Species Forum.

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Enviro-Responsible

DECEMBER 23, 2025



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Introduction

- Aquatic invasive species (AIS) threaten **ecosystems, fisheries, and infrastructure**
- Early detection is critical for **rapid response and cost control**
- Traditional microscopy:
 - Reliable for **established populations**
 - Poor sensitivity for **early-stage invasions**
- eDNA offers a **high-sensitivity alternative**



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Objective

This study assessed the presence of three Aquatic Invasive Species (AIS) *Dreissena polymorpha*, *Dreissena bugensis*, and *Bythotrephes spp.* in samples collected between 2022 and 2024 from several lakes of the Manitoba basin using both conventional microscopy and environmental DNA analysis.

- **Why these species?**
- High ecological and economic impact
- Known or emerging presence in Manitoba waters





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Study Sites

- **Lake Manitoba:** Established zebra mussel populations
- **Cedar Lake:** Spiny water flea hotspot; emerging dreissenid signals





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Study Objectives

- To compare **microscopy vs. eDNA (qPCR)** detection
- To assess **sensitivity and agreement** between methods
- To identify **early-stage invasions** missed by microscopy because of very low-density populations
- To better inform **integrated AIS monitoring strategies**

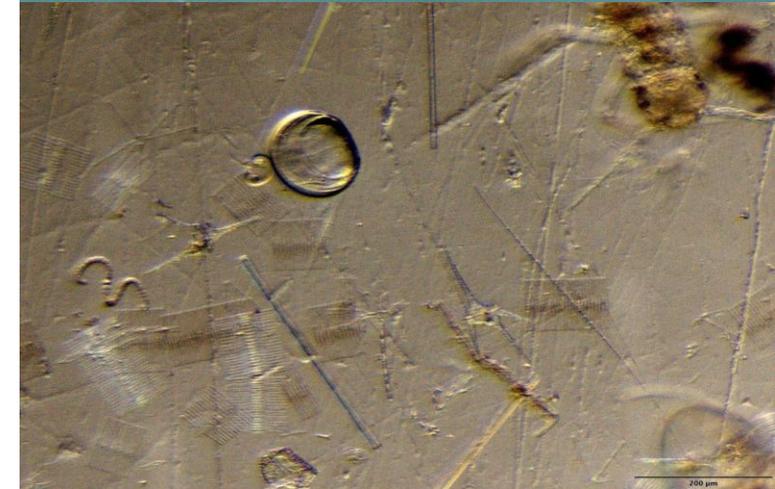
Sampling overview

- **29 water samples (70% ethanol-preserved)**
- **Collected across 21 sites (2022–2024)**
- **Parallel analysis using:**
- **Light microscopy and polarized light filter for veligers**
- **eDNA extraction + qPCR**



Microscopy Methodology

- Visual identification of:
 - Adults of SWF
 - Larvae (veligers)
 - Zooplankton stages
- Strengths:
 - Confirms established populations, **viable and dead organisms, as well as SWF embryos and resting eggs.**



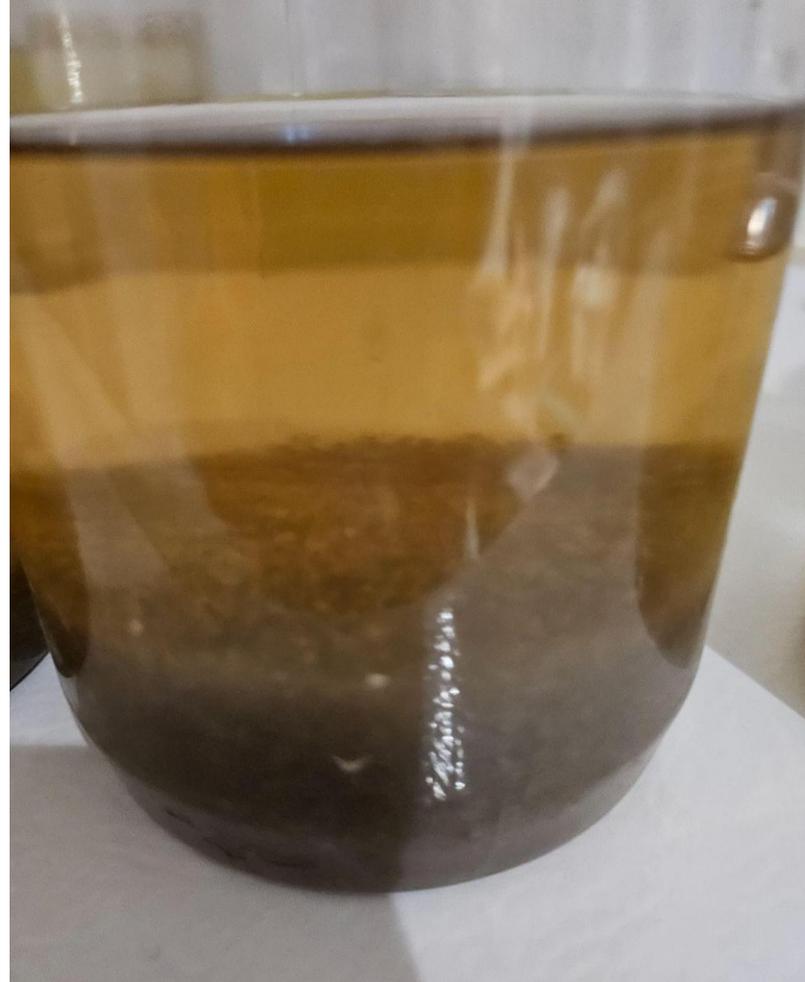


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Microscopy Methodology

- Limitations:
 - Low sensitivity at very **low abundance of organisms or early invasion**
 - Time consuming for samples with extremely high density plankton, including algae, and/or sediment.

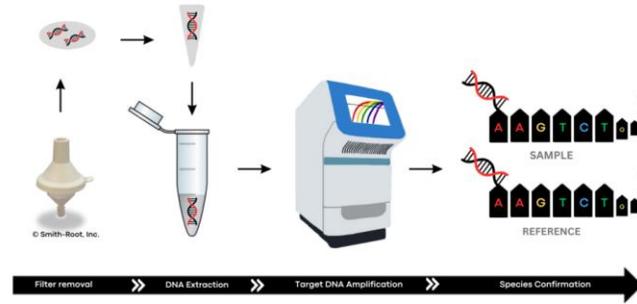


Plankton

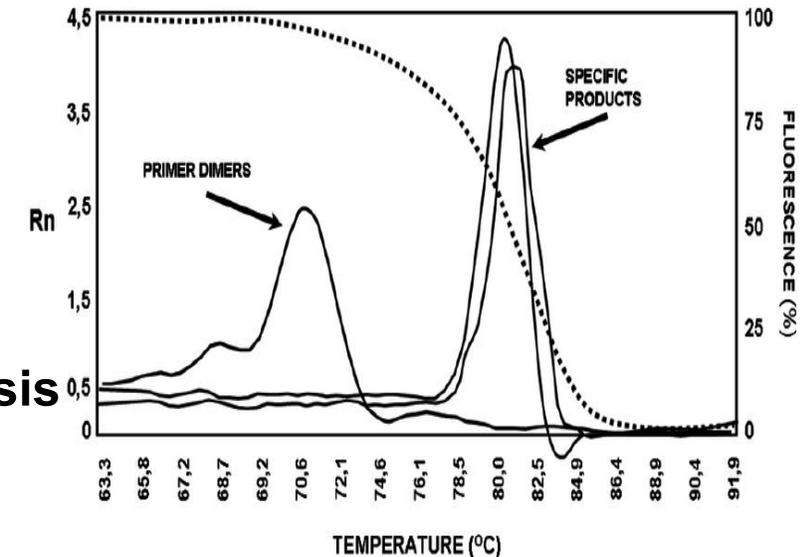
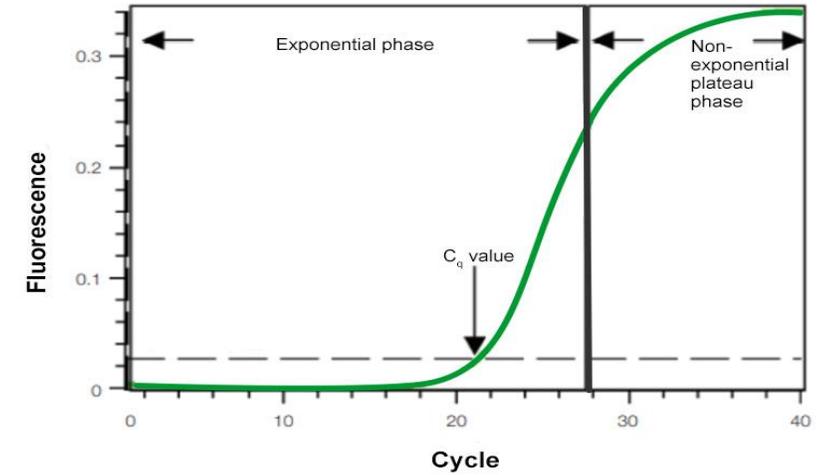
**Sediment in plankton
sample**



eDNA & qPCR Workflow



- Filtration (26 mm pore-size filter)
- DNA extraction (PowerWater kit)
- qPCR (SYBR Green):
- COI (species-specific)
- 18S (internal control)
- Positive = **Ct ≤ 40**
- Verification by **melt curve + gel electrophoresis**

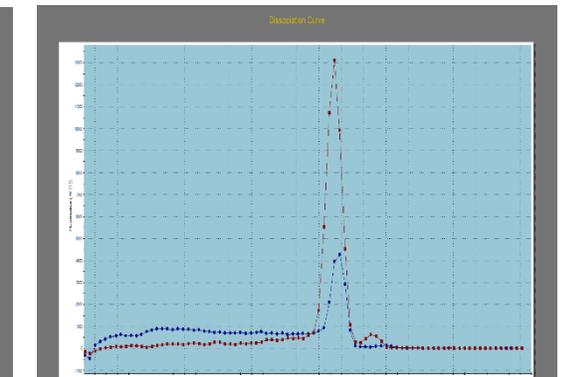
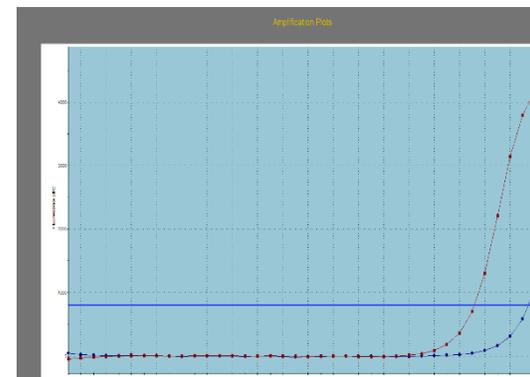
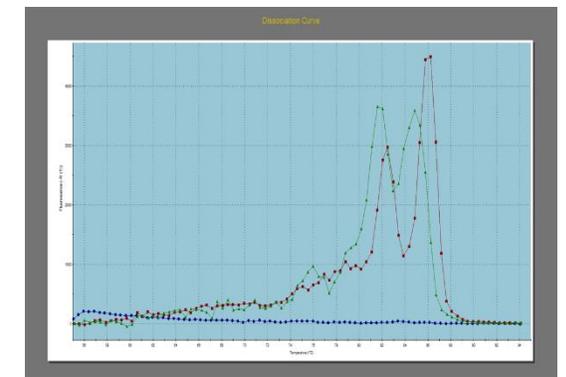
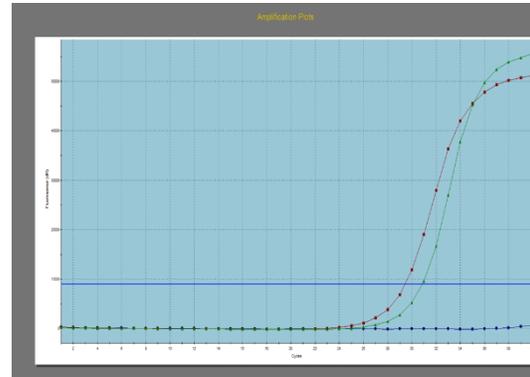
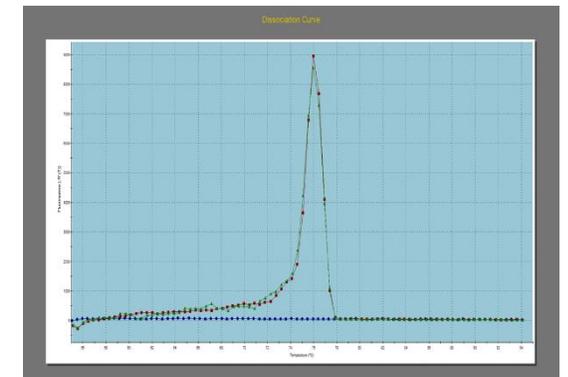
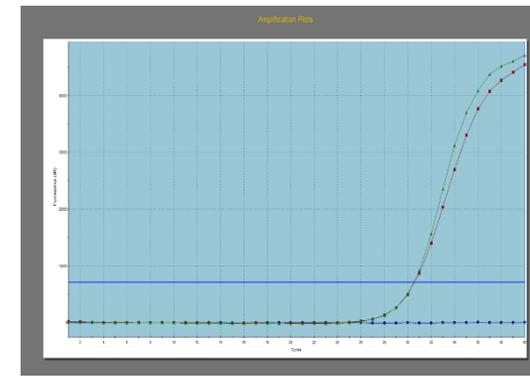


eDNA targets amplification

- Clear amplification:
- Ct range: 28–36
- Single melt peaks

Primer ID	Primer sequence	Reference
Bytho_F	GCAGGAACTGGCTGAACA	Walsh et.al., (2019)
Bytho-R	AATAATAAAAGGAGGGCTGTAATACC	
COI-DP -F3	TAATGGGGGGATTCCGGAA	Williams et al. 2017
COI-DP -B3	GCTCCCCCAATATGAAGAG	
COI-DB-F3	ATTTGGTGGGGTTGAAC	Williams et al. 2017
COI-DB--B3	GGCTAAAACAGGTATTGCTAA	
18S -F3	GTTAGCCCAGACCAACGC	Williams et al. 2017
18S -B3	CTTCCTTGGATGTGGTAGCC	

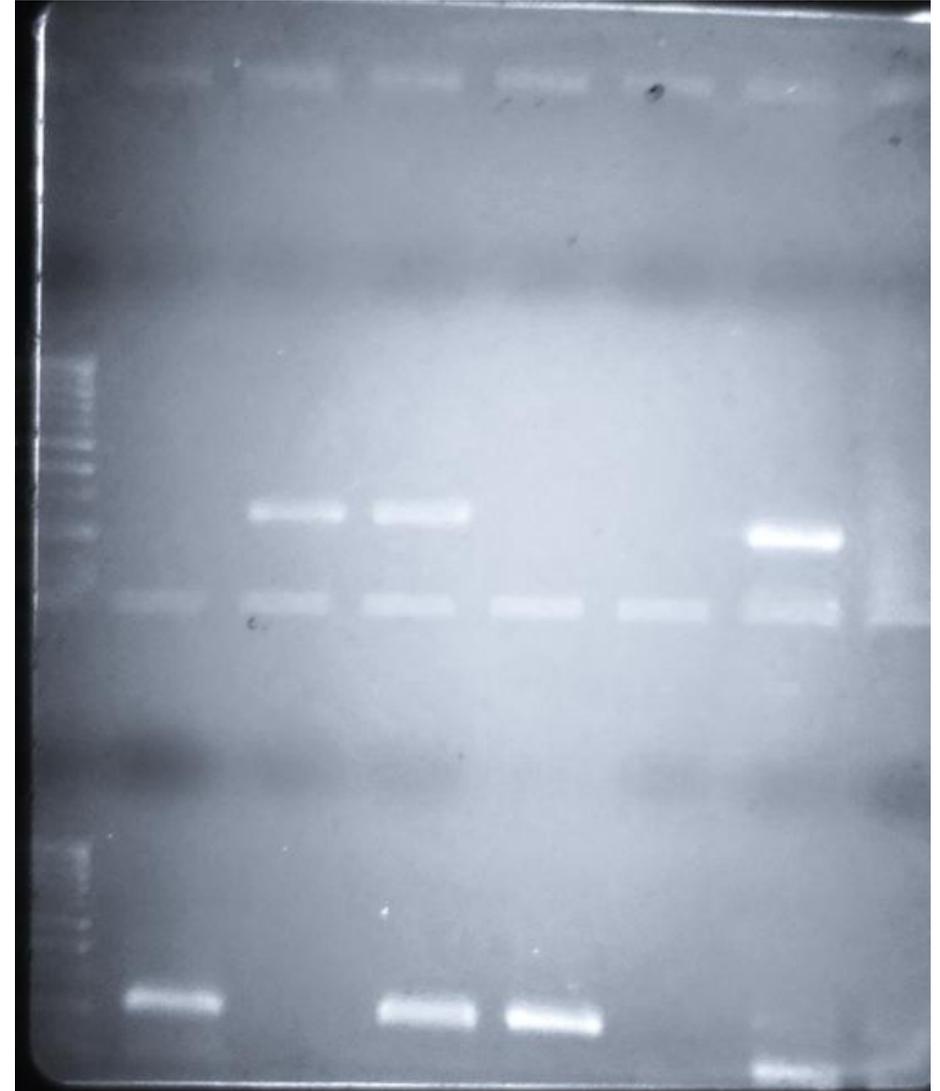
Fig 2 Amplification plots and dissociation profiles for the COI and 18s DNA segments using the primer pairs developed by William et. al, 2021. A) Amplification of *D. polymorpha* COI in eDNA samples V46 and V47 and NTC, B) Dissociation profile for the COI target in samples V46 and V47 and NTC; C) Amplification of the bivalve small subunit ribosomal RNA (18s) in eDNA samples V46 and V47 and NTC; D) Dissociation profile for the 18s target in samples V46 and V47 and NTC; E) Amplification of *Bytrothrephes longimanus* in samples V4 and V5 from Cedar Lake F) Dissociation profile for the *Bytrothrephes longimanus* target in samples V4 and V5 and NTC.



qPCR Validation Results

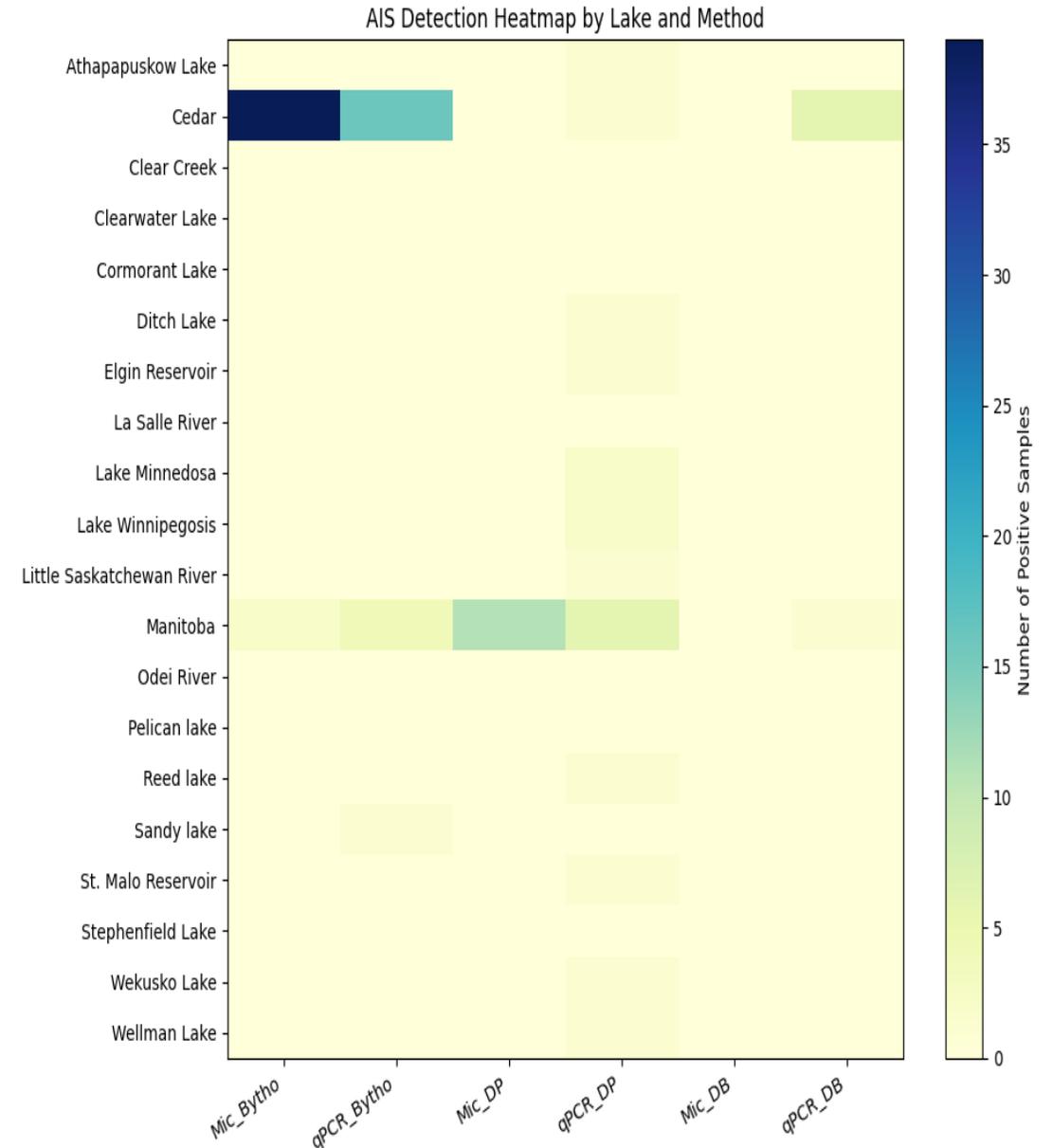
- Agarose gels:
 - Single bands (~200 bp)
 - No amplification in NTCs
- Confirms **specificity and reliability**

Fig.3 Electrophoresis of qPCR amplification products in 1.5% agarose. Single bands were observed for the target species *D polymorpha* COI and *Bythotrophes sp.* (COI) with sizes around 200 bp with no amplifications in the non-template control samples (NTC).



Key Results: Detection Patterns

- **Microscopy**
 - *Bythotrephes*: 39 detections, **Cedar Lake only**
 - *Zebra mussel*: 13 detections, **Lake Manitoba only**
- **eDNA (qPCR)**
 - *Bythotrephes*: 29 detections across **8 lakes**
 - *Zebra mussel*: 21 detections across **12 lakes**
 - *Quagga mussel*: Rare or absent
-  eDNA reveals **broader, low-density distribution**





Method Agreement (Cohen's κ)

Species	TP	FN	FP	TN	Sensitivity	Specificity	κ (Kappa)
<i>Bythotrephes sp.</i>	15	26	7	55	0.37	0.89	0.27
<i>Dreissena polymorpha</i>	3	8	18	74	0.27	0.80	0.06
<i>Dreissena bugensis</i>	0	0	7	96	NA	0.93	0.00

- TP (True Positive)**: detected by both microscopy and qPCR
- FN (False Negative)**: detected by microscopy, missed by qPCR
- FP (False Positive)**: missed by microscopy, detected by qPCR
- TN (True Negative)**: not detected by either method

Agreement between microscopy and qPCR was low to fair across taxa ($\kappa = 0.00$ – 0.27). Sensitivity of qPCR relative to microscopy ranged from 0.27 to 0.37, while specificity remained high (0.80–0.93). For *Dreissena bugensis*, no microscopy detections were recorded, resulting in zero agreement despite seven qPCR positives.

Why Methods Disagree

- DNA degradation in stored samples
- PCR inhibition
- Patchy organism distribution
- Microscopy detects **presence and density of established populations**
- eDNA detects **genetic signal**, not viability
-  Differences are **expected and informative**



Lake-Specific Insights

- **Lake Manitoba**
 - Zebra mussels eDNA dominant and persistent
 - Quagga mussels eDNA signal rare
 - Patchy spiny water flea signals
- **Cedar Lake**
 - Strong eDNA *Bythotrephes* establishment
 - Quagga mussel eDNA signal expansion suggested
 - Zebra mussel eDNA confirmed in **veligers (2025)**



*The language used in this slide has been changed from the original presented at the FORUM to avoid confusion.



Management Implications

Aspect	Microscopy	qPCR
Sensitivity	Low–moderate	High
Early detection	✗ Poor	✓ Strong
Confirms live organisms	✓ Yes	✗ DNA only
Best use case	Established populations	Early invasion surveillance

Conclusions



**eDNA analysis
outperforms
microscopy for
early detection**



Microscopy remains
critical for
confirmation of
established
populations.

=
EQUALS

**Combined approach
= Best practice**



Supports proactive AIS
management in Manitoba
and in Canada



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Acknowledgements

- A very special thank you to Joey Tonin, Aquatic Biologist, and others at the Aquatic Invasive Species Unit (AISU), Department of Environment and Climate Change in Manitoba for their support in providing the samples and allowing us to use the data for the eDNA analysis. Thank you for your support over the last few years and for the collaboration with this project.
- A big thank you to the Invasive Species Centre for the opportunity to participate and share our research in this FORUM, and for organizing this and many other events.

Thank you everyone!

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