# **Environmental DNA (eDNA)**

What eDNA can do for you, and how you can begin utilizing this powerful tool within your watershed











James Garner - PhD Student at UMass Amherst Co-advised by: Michelle Staudinger and Adrian Jordaan











Photo Credit: Jimmy Powell

## My Background

• Previously worked with the Massachusetts Division of Marine Fisheries (MA DMF) as a Biological Fisheries Technician



## My Background



 1<sup>st</sup> year of PhD – I worked part time as the Ecology Program Director for the Jones River Watershed Association (where eDNA projects began)



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- examples: soil
  - (Ancient DNA)







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- examples: soil, water, air, honey!



Ribani *et al*. 2020

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- examples: soil, water, air, honey!
- Like a genetic NOSE for a given environment



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  - Relative abundance\*

\* Technique for calculating relative abundance is still being refined and requires calibration to other established techniques

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Photo Credit: Jimmy Powell – JRWA SmugMug

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Estimates of species density/abundance	Provide a low-cost supplement or alternative to other methods

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- This is a perfect step to get communities and citizen scientists involved with a project!



Aman, J., Kinnison PhD, M. T., Holmes, V., & Gottsegen, C. (2020). Developing Cost Effective Monitoring for Rainbow Smelt Using eDNA.



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- Filtration



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- Analysis



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For both approaches, the first THREE steps are (basically) the same



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1	METABARCODING DATA						
2	Taxon	sb	lr	hb	1b	2b	3b
4	American eel	1	1	1	1	0	1
5	american shad	0	0	0	0	0	0
6	Atlantic tomcod	0	0	0	0	0	0
7	blueback herring	0	1	0	0	0	1
8	brook trout	0	0	0	1	0	1
9	hickory shad	0	0	0	0	0	0
10	rainbow smelt	0	1	0	0	0	0
11	sea lamprey	0	0	0	0	0	1
12	striped sea-bass	0	0	0	0	0	0
13	white perch	0	0	0	0	0	0
14	Atlantic herring	0	0	0	0	0	0
15	Atlantic silverside	0	0	0	0	0	0
16	fourspine stickleback	0	0	0	0	0	0
17	grubby sculpin	0	0	0	0	0	0
8	haddock	0	0	0	0	0	0
19	menhaden	0	0	0	0	0	0
20	mummichog	0	0	1	0	0	0
21	ninespine stickleback	0	0	0	0	0	0
22	rock gunnel	0	0	0	0	0	0
23	sheepshead minnow	0	0	1	0	0	0
24	striped killifish	0	0	0	0	0	0
25	winter flounder	0	0	0	0	0	0
26	black crappie	0	0	0	0	0	0
27	black crappie or rock bass	1	1	1	1	0	1
28	bluegill sunfish	1	1	1	1	0	1
29	brown bullhead	0	1	0	0	0	0
30	chain pickerel	1	1	1	1	0	1



# What can eDNA tell us?

- Species presence
- Biodiversity
- Estimates of abundance (calibrated)

# What can eDNA tell us?

- Consider what your data is telling you compared to other traditional monitoring techniques
  - What are some other traditional ways we monitor aquatic life?

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# Things eDNA can't tell you







# Things eDNA can't tell you

• Absolute abundance









# Things eDNA can't tell you

- Absolute abundance
- Age and Growth structure for fish populations







# Things eDNA can't tell you

- Absolute abundance
- Age and Growth structure for fish populations
- What life stage your DNA signal came from (yet)







eDNA is a SUPPLEMENT to ongoing monitoring practices, not a replacement.

## Strengths/Weaknesses

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## Pros:

- Cost effective
- Less upfront effort
- More accessible than traditional ecological monitoring
- Non-invasive/destructive
- Rare, shy, and cryptic species detection
- Field constantly being refined
- Novel applications emerging regularly

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## Cons:

- False positives/negatives common
- Controversial (abundance estimates)
- Barrier to entry for downstream analyses (after sample collection/filtration) extremely high
- Still a new field requiring refinement

## Known biases

# eDNA monitoring techniques have limits and sampling biases.





## The importance of calibration



Figure SF8. Correlation between species richness observations from conventional surveys and eDNA metabarcoding for freshwater (red, n=104) and marine systems (blue, n=17). The line represents a 1:1 relationship.



#### Chambert et al. 2017

#### McElroy et al. 2020

## One more time...

eDNA monitoring techniques are not meant to REPLACE other monitoring strategies, but to ENHANCE ongoing monitoring efforts

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or

Provide an accessible and affordable starting point for watershed, biodiversity, and species monitoring efforts

# eDNA monitoring is a "Force Multiplier" for other monitoring efforts





Determining IF your restoration/adaptation action met its intended goals

- Pre/post dam removal monitoring through time
- Pre/post fishway installation monitoring, etc.

Diadromous Fish Species Diversity Pre and Post Mainstem Dam Removal



Determining how far upstream river herring are making it into your watershed



## Whether or not river herring are making it upstream of fish ladders



https://www.nsrwa.org/protect-our-waters/healthy-rivers/dam-removals/south-river-restoration/

When EXACTLY river herring are arriving to your watershed, how long they stay, and when they have all left (sampling through time)

Robbins Pond	
Snipatuit Pond	
Long Pond	
Billington Sea	
Furnace Pond	
Oldham Pond	
Great Herring Pond	
Cedar Lake	
Pilgrim Lake	
Johns Pond	
Santuit Pond	
Lower Millpond Pond	
Upper Mill/Walkers Pond	
Gull Pond	
Coonamessett Pond	
Whitmans Pond	
Chebacco Lake	
Pentucket Pond	
Upper Mystic Lake	
Lower Mystic Lake	
12 - 121 - 12	1 . 14 . 192 . 193 . 193 . 193 . 199 . 199 . 19 . 19



Rosset, J., Roy, A. H., Gahagan, B. I., Whiteley, A. R., Armstrong, M. P., Sheppard, J. J., & Jordaan, A. (2017). Temporal patterns of nigration and spawning of river herring in coastal Massachusetts. *Transactions of the American Fisheries Society*, *146*(6), 1101-1114.

If river herring are entering your watershed from other locations

• other inlet streams to a headwater pond, for example



Biodiversity and community structure of:

- Prey species
- Habitat/plant community structure
- Aquatic pathogens
- Freshwater competitors





Provide low-cost biological evidence for permitting



## Things to consider: Implementing eDNA in your watershed

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- Time series/sampling locations

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- Ship frozen water samples or preserved (filtered) samples to a lab

#### Some New England based Labs you can work with:

#### **University of Maine, Kinnison Lab**

https://umaine.edu/evoappslab/people/dr-michael-kinnison/

UMaine eDNA website: https://umaine.edu/edna/

Contact: Geneva York: <u>geneva.york@maine.edu</u>

#### **University of New Hampshire, Hubbard Center for Genome Studies**

https://hcgs.unh.edu

Contacts: Krystalynne Morris: <u>krystalynne.morris@unh.edu</u> Kelley Thomas: <u>kelley.thomas@unh.edu</u>

## What's it going to cost me?

(shipping and handling not included)

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Biodiversity (Metabarcoding): Roughly between \$2,000 and \$3,000 for 96 samples

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> Single Species (qPCR): Less than \$20 per sample



### Some of my ongoing work



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The Goal of my PhD research is to make eDNA tools and techniques more accessible, and to get them in the hands of the people, communities, and managers who can use them the most.



## Some of my ongoing work

 Project 1: Using eDNA biodiversity methods to measure restoration action efficacy in a coastal New England watershed

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encing, filtration, comparing novel eDNA biodiversity encing, filtration, and preservation techniques to https://www.letd.deployable.biodiversity filtration.

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Project 2: Calibrating eDNA abundance metrics to established methods (electronic herring counters, fyke net surveys, purse seine surveys, and electrofishing surveys)









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#### Project 4:

Developing a field deployable, affordable, equitable, and accessible eDNA biodiversity monitoring kit

Project being tested/validated in conjunction with the Town of Plymouth, MA in preparation for an upcoming dam bypass project



microbial and restoration

estoration monitoring





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- Project 4: Validating/comparing novel eDNA biodiversity sequencing, filtration, and preservation techniques to develop a turn-key, field deployable biodiversity monitoring system

### Resources and Literature

- Aman, J., Kinnison PhD, M. T., Holmes, V., & Gottsegen, C. (2020). Developing Cost Effective Monitoring for Rainbow Smelt Using eDNA.
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Thank you!!!

For references, contacts, or any questions, please don't hesitate to reach out!

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