### The use of eDNA in comparison to trawl and historical catch data for assessing marine finfish in Matagorda Bay, Texas

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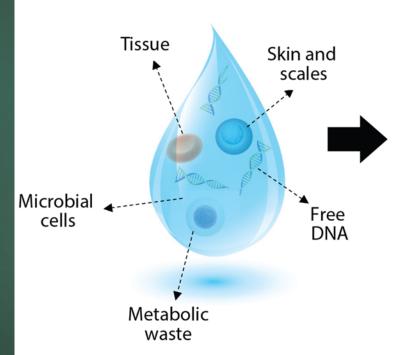
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### Environmental DNA (eDNA)

An emerging tool for studies of presence and abundance of aquatic organisms

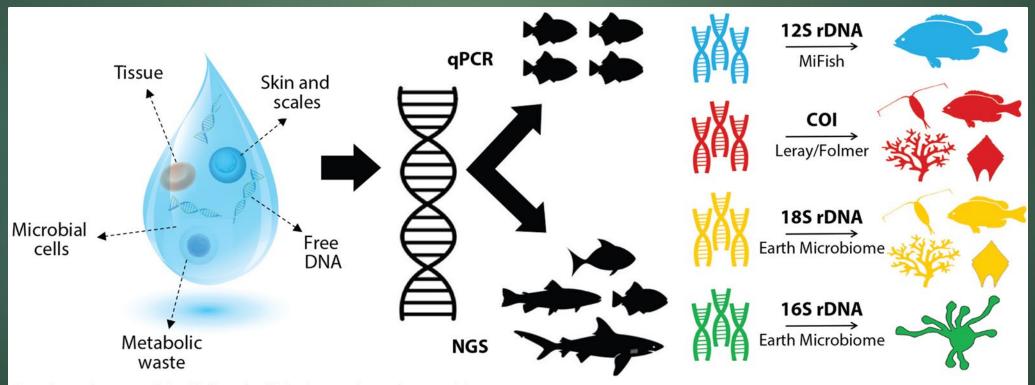
minimally invasive methodology (water sampling)



https://tos.org/oceanography/article/observing-life-in-the-sea-using-environmental-dna

## Environmental DNA (eDNA)

- flexibility to accurately detect either
  - Single species specific
  - multi-species biodiversity in marine communities.
- Being deployed in the context of management worldwide



https://tos.org/oceanography/article/observing-life-in-the-sea-using-environmental-dna

## Texas Parks & Wildlife (TPWD)

- Texas Parks and Wildlife Department's (TPWD) extensive independent fisheries monitoring program
- Several sampling techniques that provides a comprehensive means of assessing finfish communities
  - ► Gill Nets
  - Bag Seines
  - Shrimp Trawls (Trawls)





### Objective

Compare the fish community of Matagorda Bay as inferred by eDNA metabarcoding with a 47-year dataset of fisheryindependent sampling conducted by TPWD in Matagorda Bay using bag seines, trawls, and gill nets.







### Methods – Field Sampling 2023

#### Matagorda Bay

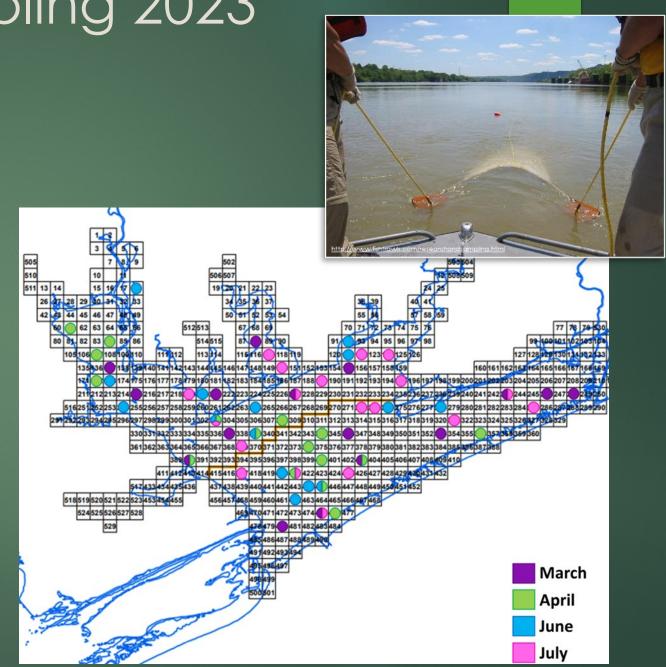
- Randomized sampling
- Aimed for 20 per month

#### eDNA

- 2 surface water samples per trawl
- Taken before trawl was used
- Placed on ice

#### Trawls

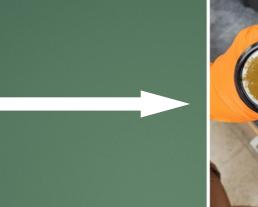
- Minimum depth of 1 meter
- ► 10-minute tows
- Towed in circular pattern
- Worked up in field
- Historical data from TPWD's database
  - Gear bag seines, trawls, and gill nets
  - ► Time from 1975-2022 for the 4 months



### Methods – Lab Processing

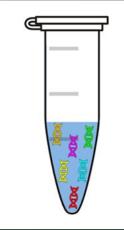


Filtration Peristaltic Pump & Smith-Root eDNA glass filter pack





DNA Isolation Qiagen PowerSoil Pro Kit & Zymo PCR Inhibitor Removal Kit



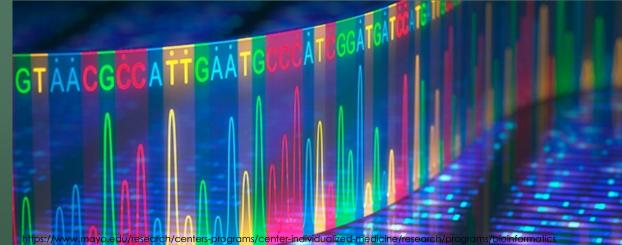
Library Prep Metabarcoding (MiFish primers)



Illumina Sequencing (Texas A&M Agrilife)

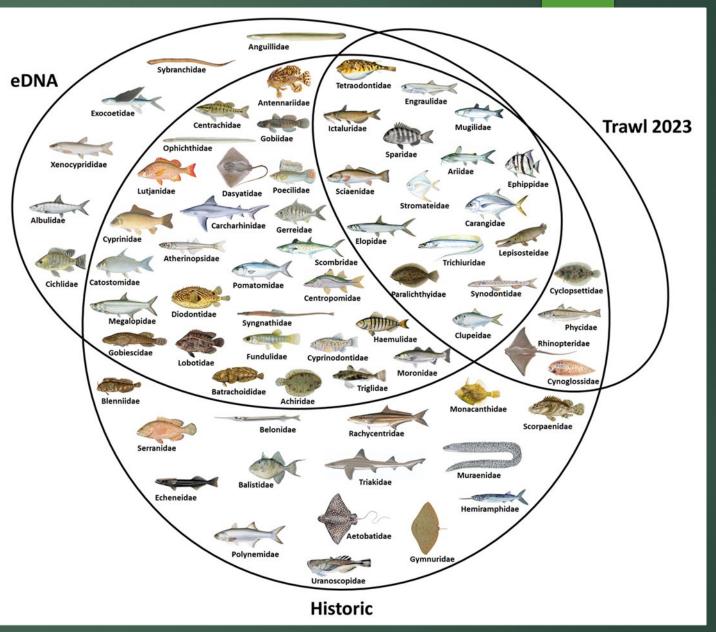
### Methods – Analysis

- Sequences were Demultiplexed, trimmed, and compared against reference library using DADA2 bioinformatics tools
- Compared presence/absence among traditional gears and eDNA
  - Comparisons made at family and species level
  - Comparisons gear relationships in the 2023 trawls and eDNA
    - Top 10 species caught and detected
    - Monthly species accumulation curves
- Compared species richness among traditional gears and eDNA using nonparametric Wilcoxon



### Results – Family Level Comparisons

- Historically, 62 families have been detected between 1975-2022
- Trawl 2023, detected 20 families
- eDNA
  - 16 (80%) of the 2023 trawl detected families
  - 49 (79%) of the historically detected families
  - eDNA detect 6 additional families not detected with traditional methods.



### Results – Species Level Comparisons

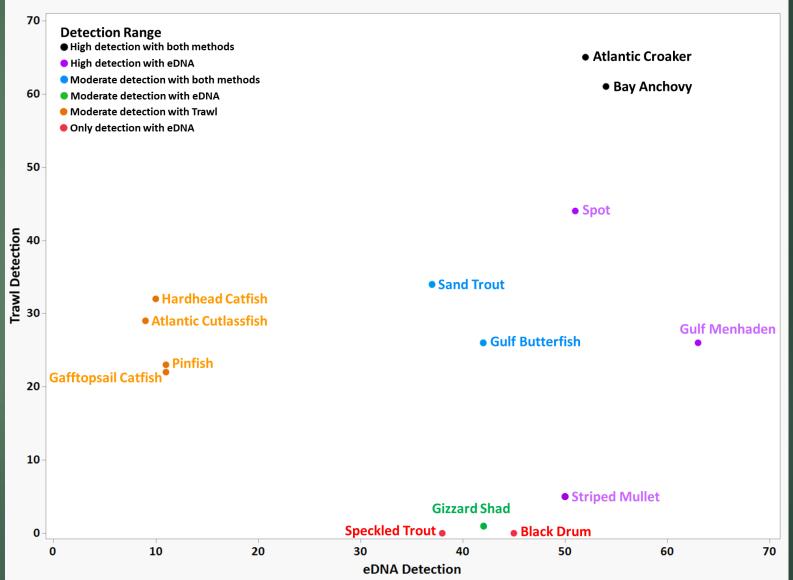
#### General

- ► Historical = 168
- ► Trawl = 36
- ▶ eDNA = 105
- The top 3 species for the historical and trawl catch were present in the top 5 species detected in the eDNA

Gear	#1	# 2	#3	# 4	# 5
Historical Catch (1975-2022)					
	Gulf Menhaden (32%)	Atlantic Croaker (20%)	Spot (11%)	Hardhead Catfish (6%)	Pinfish (3%)
Trawl Catch (2023)					
	Atlantic Croaker (71%)	Bay Anchovy (16%)	Spot (5%)	Sand Trout (3%)	Pinfish (1%)
eDNA Detection (2023)					
	Gulf Menhaden (94%)	Bay Anchovy (81%)	Atlantic Croaker (78%)	Spot (76%)	Striped Mullet (75%)

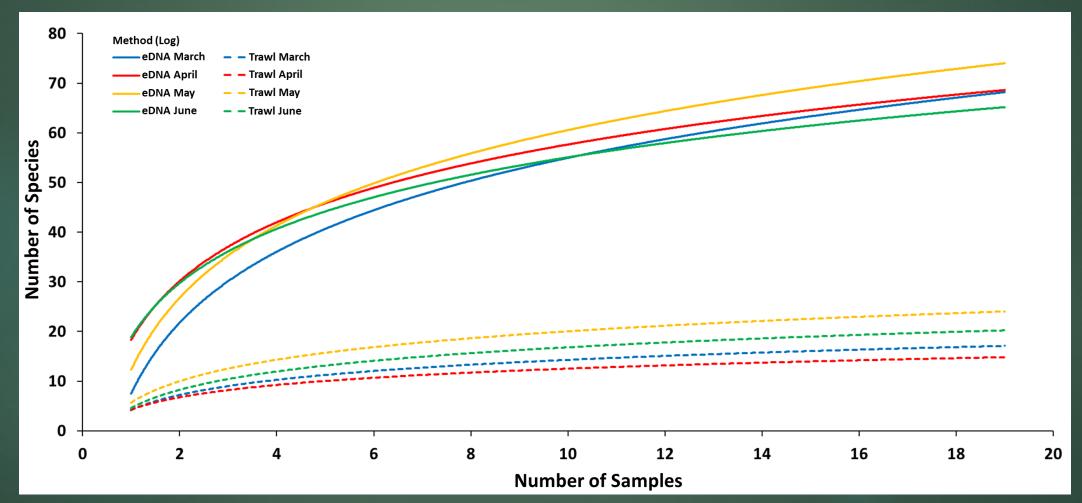
### Results – Detection Range Comparisons

- Top 10 species detected in both methods
- Atlantic croaker and bay anchovy are highly detectable by both methods
- Other species like gulf menhaden or hardhead catfish were detected better by either trawls or eDNA
- Speckled trout and black drum were ONLY detected in eDNA



### Results – Species Accumulation Curves

Validated the finding that higher levels of species detection could be expected with eDNA than trawls

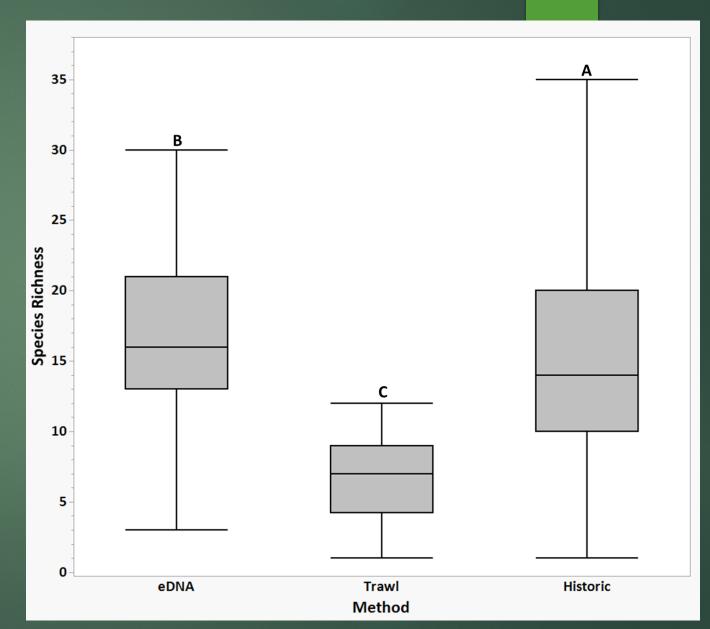


### Results – Species Richness

► Kruskal – Wallis = <0.001

#### ► Wilcoxon

- Highly significantly (<0.001) between eDNA and 2023 trawls, and historical data and 2023 trawls
- Slightly significant (0.033) between eDNA and historical



### Discussion

- Overall eDNA detected over 79% of the families present in the historical record
- Top 4 present species in eDNA were in the top 3 species for both the trawl and historical catches
- Side-by-side sampling (eDNA vs trawl) showed that eDNA could detect a greater diversity of fish species than trawls. This could be caused by...



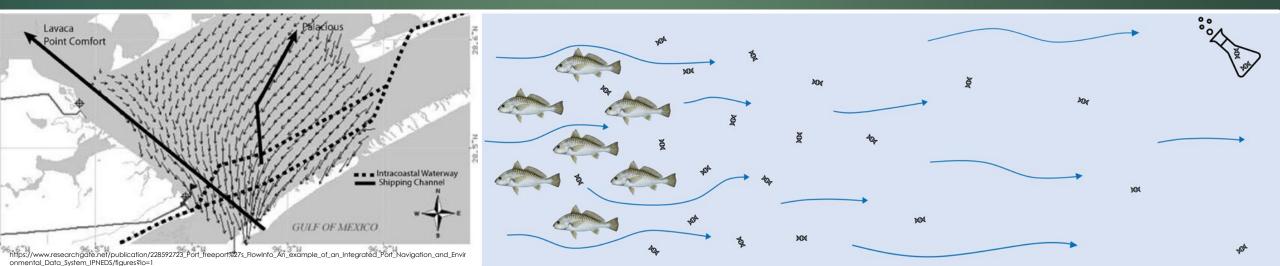
### Discussion

- Gear bias size-selection of the trawls, may prevent the detection of small- and large-bodied species
- Trawl deployment occurs in deeper water away from structures (e.g. reefs and jetties) preventing sampling shallow-water and structure-associated species
- TPWD trawl sampling occurs during daylight hours, decreasing the chances of detecting nocturnal species that spend the day resting in burrows or dens.



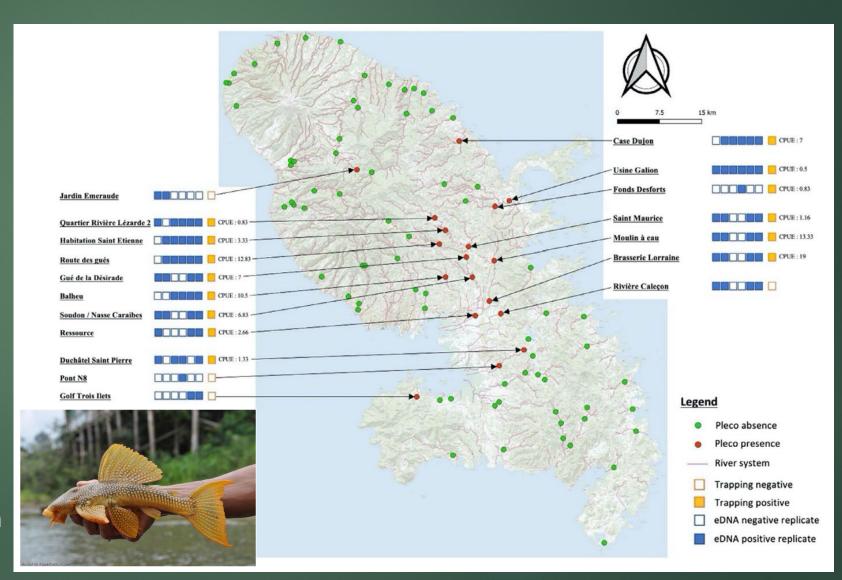
### Discussion

- eDNA drifts in the water column
- Influenced by currents and tidal direction
- May not have been in the immediate vicinity of the trawl when the water sample was taken



## Species Specific (qPCR) Example

- ► Teta catfish
  - Armored catfish
  - Native to Trinidad and Tobago Island
- eDNA and traps set at 83 sites in Martinique Island
  - eDNA detected in 18 sites
    - Red dots = + eDNA
    - ► Green dots = eDNA
  - Traps detected in 14 site and the sites were detected by eDNA
- https://onlinelibrary.wiley. com/doi/pdf/10.1002/edn 3.260



### Acknowledgements

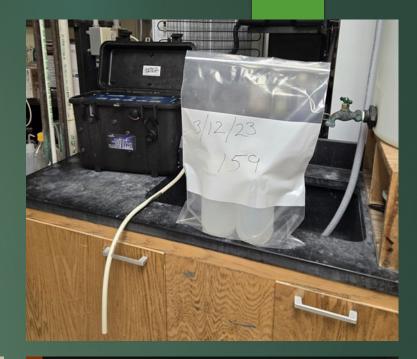
- A big thanks to my fellow colleagues at PRB and the Matagorda Ecosystem team who helped with the collections and extractions.
- Funded by a Gulf States Marine Fisheries Commission's Interjurisdictional Fisheries Program.
- Contact info
  - Polly Hajovsky
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### **Contamination Prevention**

#### Out in the Field

- Designated eDNA
  - Ice chest
  - Spot on the boat
- Nalgene bottles in Ziplock bags
- Hand sanitizer
- In Lab
  - Sterilized filter packs
  - Clean work bench and water pump with10% bleach
  - Autoclave (Nalgene bottles, forceps, etc.)
  - Set all lab materials (pipettes, tips, etc.) under a UV light
  - Designated eDNA equipment
  - Filter pipette tips







### Library Preparation Controls

- Fin clip DNA controls (n = 2)
- $\blacktriangleright$  Milliq water controls (n = 10)
- $\blacktriangleright$  Plain filter control (n = 1)

