# ASSESSMENT OF FISH SPECIES USING ENVIRONMENTAL DNA METABARCODING AND CATCH SURVEY IN THE SATELLITE LAKES OF LAKE VICTORIA, KENYA

## Joab Jared Ochingo<sup>1</sup>, Paul Meulenbroek<sup>2</sup>, Geoffrey Ong'ondo<sup>3</sup>

<sup>1</sup>Egerton University, Kenya, <sup>2</sup>University of Natural Resources and Life Sciences (BOKU), Austria, <sup>3</sup>Institute of Water Education (IHE-DELFT), Netherlands

## Background

## 1. Introduction (or educational background)

- Freshwater systems host high biodiversity but are under pressure from pollution, overfishing, habitat degradation, and climate change.
- Satellite lakes (Kanyaboli, Namboyo, Sare, Bob) are part of the Yala Wetland ecosystem and are key fish habitats.
- Traditional fish survey methods are labor-intensive and may miss cryptic or rare species.
- eDNA metabarcoding offers a non-invasive, highly sensitive tool for biodiversity assessment

#### 2. Problem statement

- Declining fish populations and lack of up-to-date species composition data hinder effective fisheries management.
- There is a need for efficient, non-invasive tools like eDNA metabarcoding to enhance fish biodiversity assessments.



Fig. 1 Anthropogenic activities in L. Kanyaboli

## **Current Work**

## 1. Specific Objectives

- To determine the fish species composition in Lakes Bob, Kanyaboli, Namboyo and Sare using fish Catch surveys.
- To evaluate the fish species composition in Lakes Bob, Kanyaboli, Namboyo and Sare using environmental DNA metabarcoding.
- To analyze the morphometric characteristics of the fish species in Lakes Bob, Kanyaboli, Namboyo and Sare.

#### 2. Methodology

## eDNA Sampling

- 1. Water volumes filtered: Kanyaboli (3.2L), Namboyo (4L), Bob (4.7L), Sare (8.7L)
- 2. Filters preserved in buffer, followed by DNA extraction, PCR amplification, and sequencing- using Teleo primers-12SrRNA gene region
- 3. Bioinformatic analysis to identify taxa from reads
- 4. Length-weight relationship and condition factor analysed
- 5. Sorensen similarity index used to compare datasets

## 3. Results

- eDNA detected 31 taxa while catch survey found 20 taxa
- Detected O. niloticus, esculentus, variabilis, leucostictus, C. gariepinus, P aethiopicus, Pundamilia nyererei, Pundamilia pundamilia, A. argentus, H. maxillaris, astatotilapia nubilus,
- Invasive species detected (e.g., Gambusia affinis and Lates) niloticus-Nile perch ).
- Sorensen similarity index = 0.57
- LWR of O.niloticus in Kanyaboli showed an isometric growth pattern

Fig. 3 fish diversity indices.

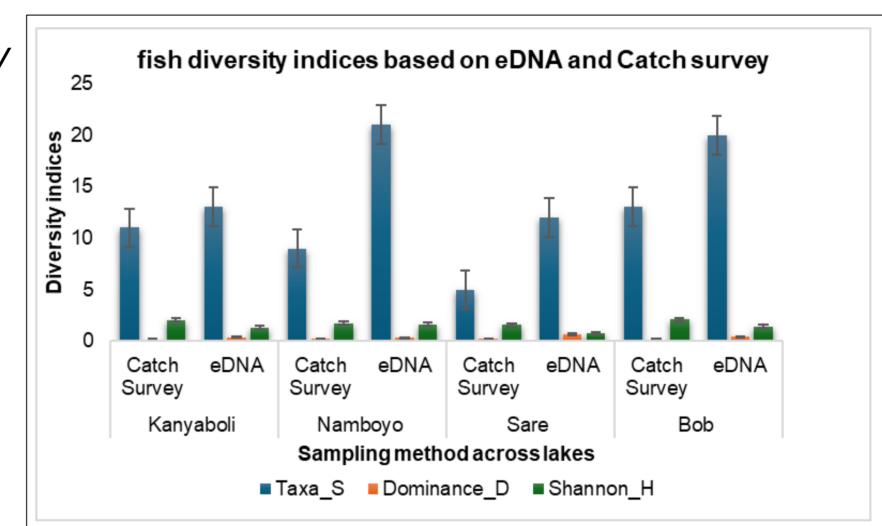
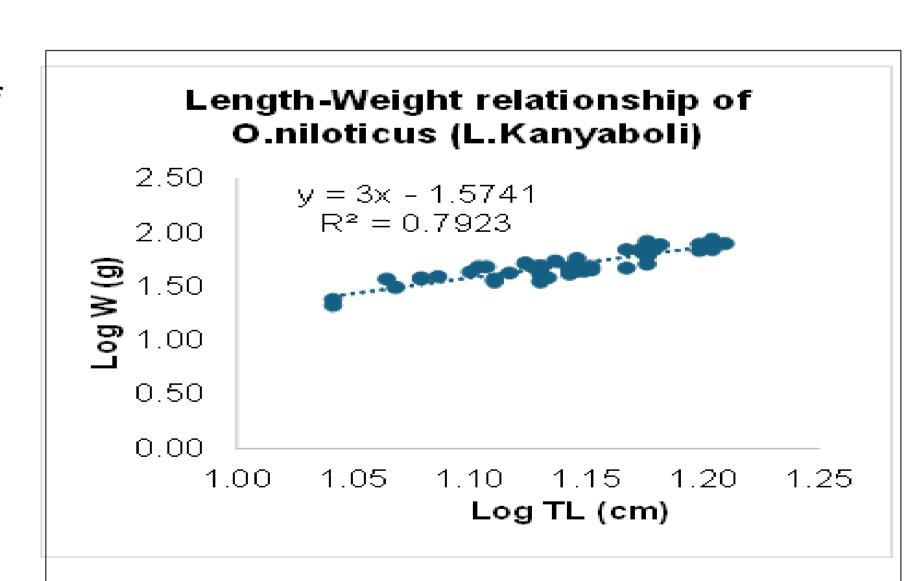


Fig. 4 LWR of O.niloticus





# **Conclusion & Expectations**

- The eDNA metabarcoding detected more fish taxa, including rare and invasive species, complementing traditional catch survey data.
- o The integration of both methods enhances biodiversity assessment and supports better conservation and fisheries management decisions.
- o The findings will guide evidence-based management, conservation, and policy formulation to protect the ecological integrity of Yala Wetland satellite lakes.













Contact: joabjared1@gmail.com P.O. Box 536 - 20115, Egerton-Njoro