



# Analysis of Environmental Factors that Correlate with the Presence of *Leptospiraceae* in a Native Hawaiian Mariculture System

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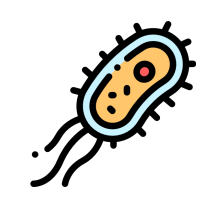
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## Introduction



Hawai'i has the **highest incidence of leptospirosis** in the United States, but cases of the disease are likely to **remain undiagnosed** or unreported.



Leptospirosis is a **zoonosis caused by *Leptospira interrogans***, transmitted via environmental dissemination through the urine of mammal carriers (rats, feral swine).



Gene *lipL32* is conservative across **pathogenic strains of *Leptospira*** and responsible for the pathogenicity of the bacteria.

## Hypothesis



We hypothesized that the **prevalence of *Leptospiraceae*** in the He'eia fishpond (HFP) positively **correlates** with significant **rainfall**, the **rainy season** of Hawai'i and **freshwater**.

## Results : Precipitation

- Leptospiraceae* presence is **significantly higher** in coastal waters during the **wet season** of Hawai'i.
- Leptospiraceae* are **strongly correlated** with precipitation during **storm** conditions.

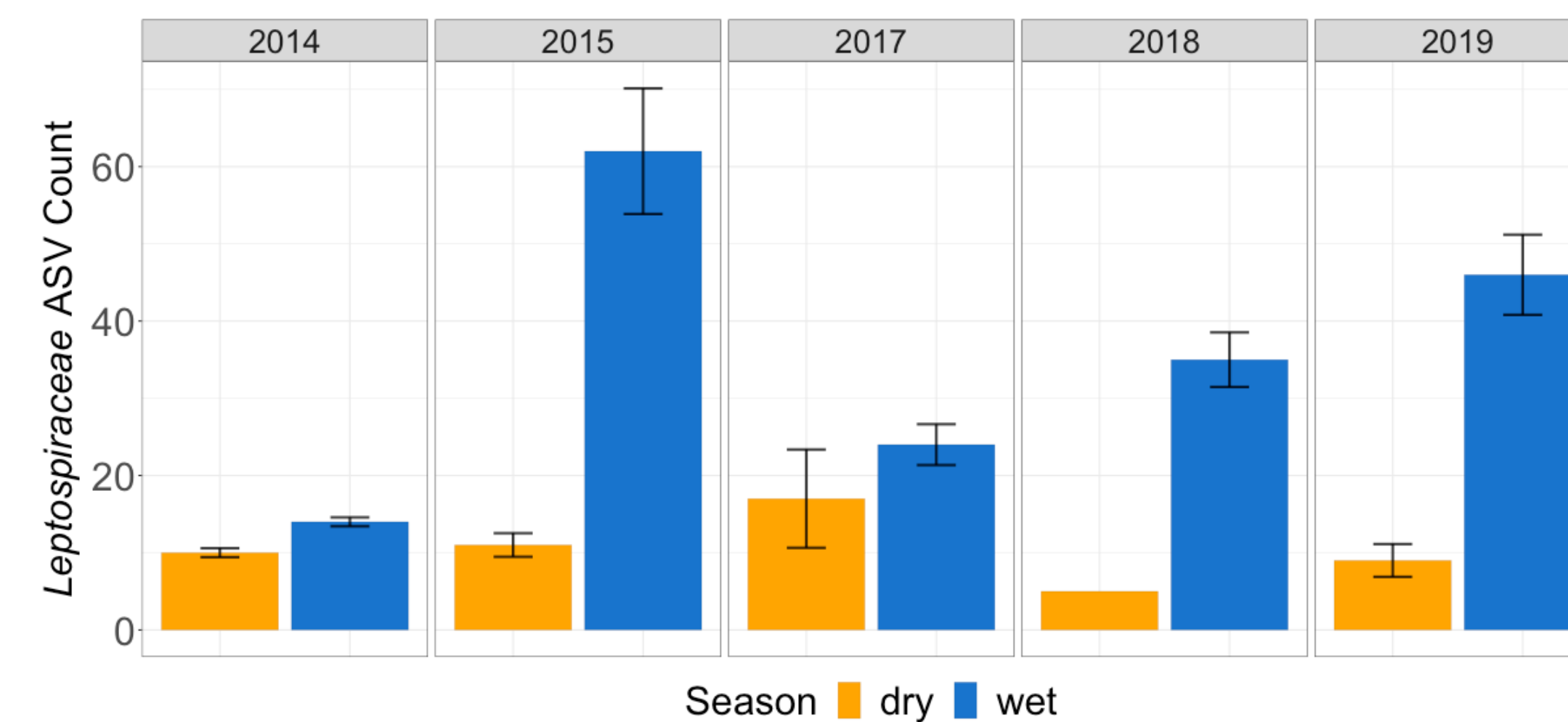


Figure 2: Comparison of the total number of *Leptospiraceae* ASV between the dry and wet seasons (2014-2015 and 2017-2019).

## Results : Salinity

- Salinity varies and total *Leptospiraceae* count vary by season.
- Leptospiraceae* is **poorly correlated** to salinity, indicating a confounding variable.

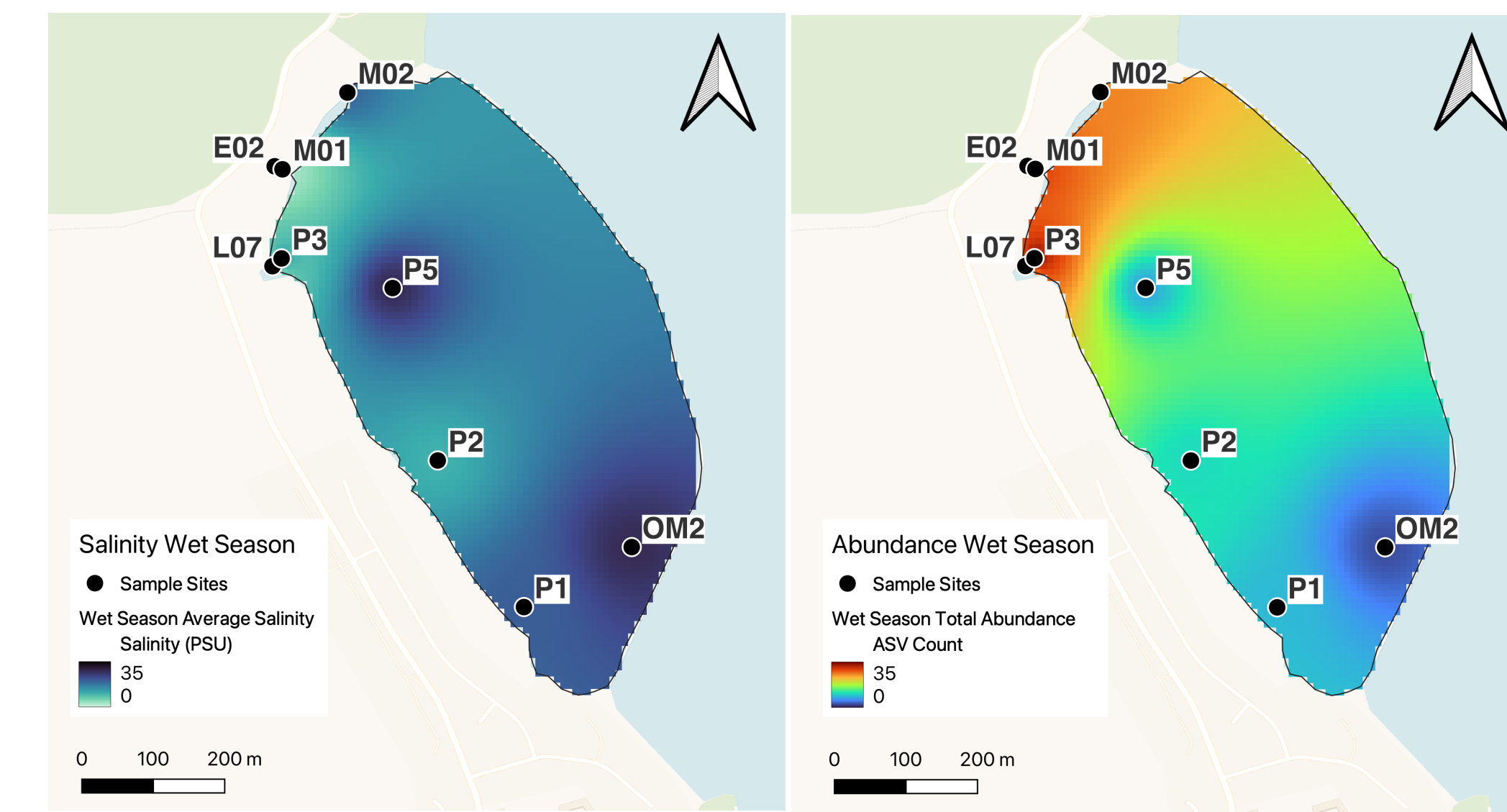


Figure 5: Average salinity (left) and total abundance of *Leptospiraceae* (right) IDW map of He'eia fishpond during the wet season.

## Riverine Input of *Leptospiraceae*

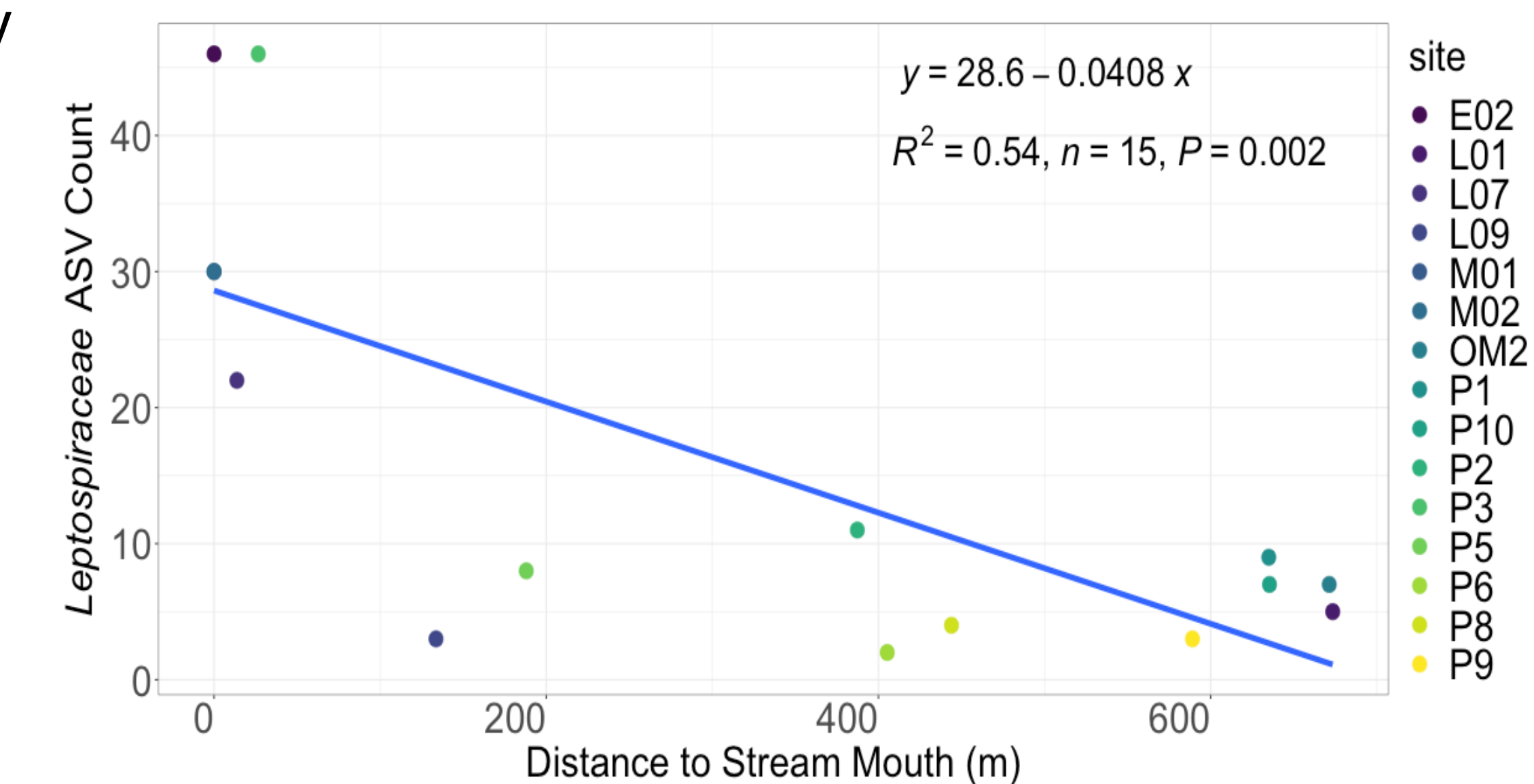


Figure 8: Good correlation between distance to stream mouth (m) and total ASV indicate a riverine input of *Leptospiraceae*.

## Discussion

- Leptospiraceae* were **significantly** more detected during the **wet season** and are **strongly associated** with **storm events**.
- Total *Leptospiraceae* ASV was **weakly correlated** with **salinity** in He'eia fishpond, but **strongly correlated** to the proximity with **He'eia stream**, suggesting a riverine input.
- This project helps to **develop methods** to understand the **circulation of *Leptospiraceae*** at the **watershed level** and inform local communities.
- Additional data** showed the **absence of pathogenic *Leptospira*** in He'eia coastal water.

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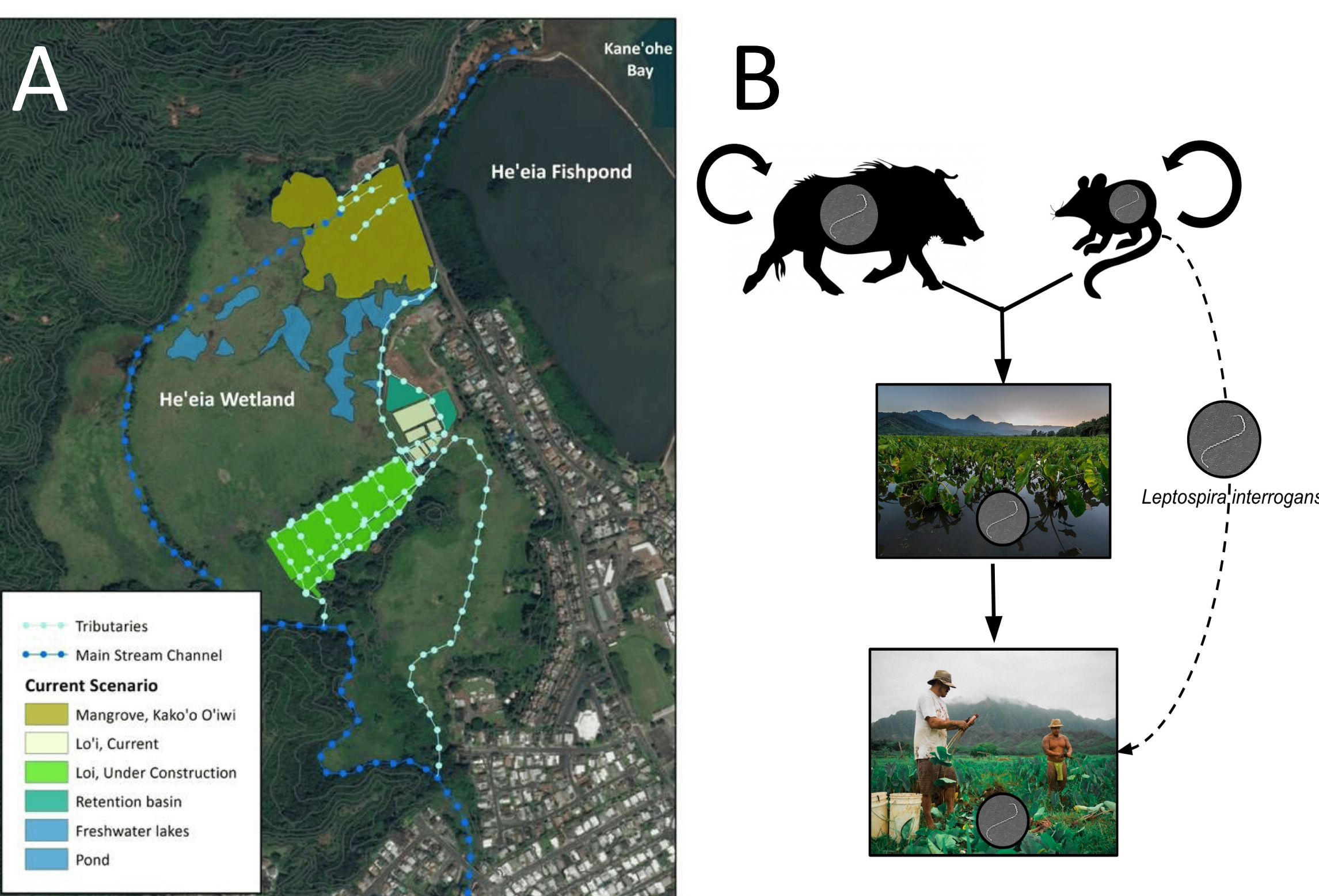


Figure 1: In He'eia watershed, traditional agro-ecological sites, channelized streams (A) and established populations of feral swine may facilitate the transmission of the disease (B).

## Methods

- Water sampling** every two weeks in 2014 - 2015 and every quarter in 2017 - 2019
- DNA extraction** and sequencing of 16S rDNA gene. Clustering of **16S amplicon sequence variants (ASV)** to identify the spatial and temporal distribution of *Leptospiraceae*.
- Salinity** data, measured using a YSI during water sampling; **precipitation** data, retrieved from the Hawai'i Climate Data Portal.
- Distance** from sample sites to **river mouth** was obtained using the measuring tool in **Google Earth Pro**.
- Data analysis and visualization performed using **RStudio** and **QGIS**.

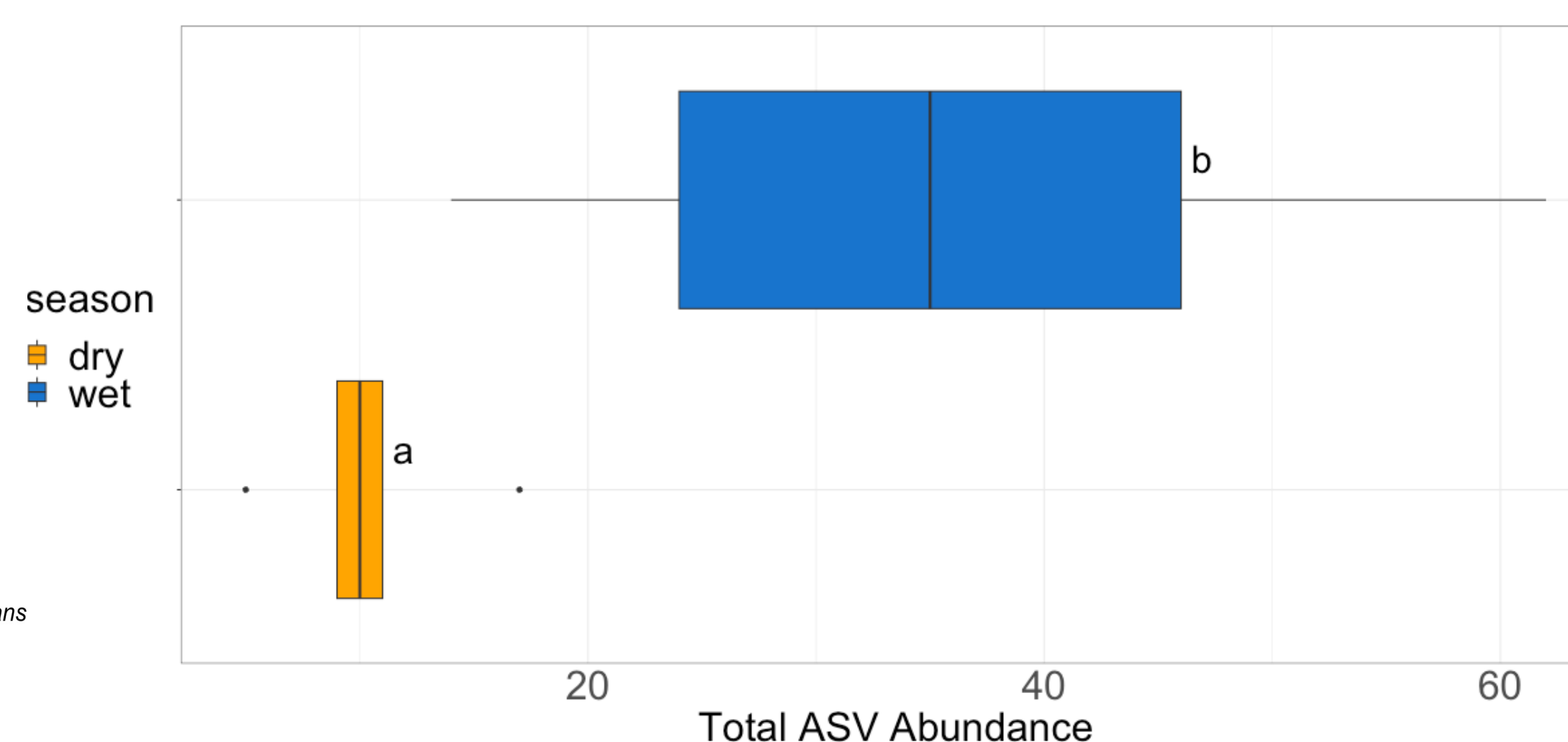


Figure 3: Variation of total ASV count between seasons (2014-2015 and 2017-2019).

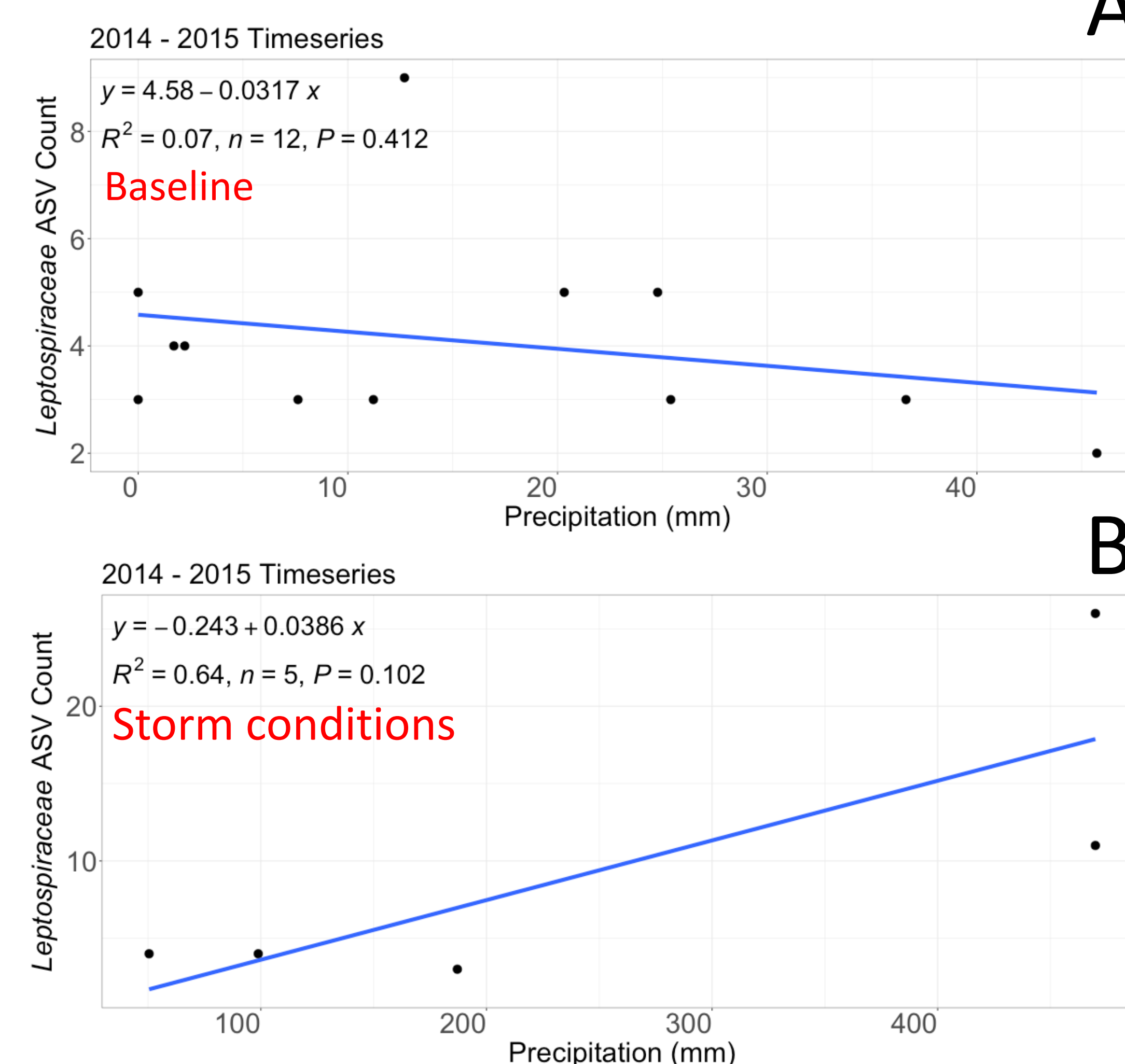


Figure 4: Correlation between *Leptospiraceae* ASV and precipitation in non-storm (A) and storm (B) conditions.

Figure 6: Average salinity (left) and total abundance of *Leptospiraceae* (right) IDW map of He'eia fishpond during the dry season.

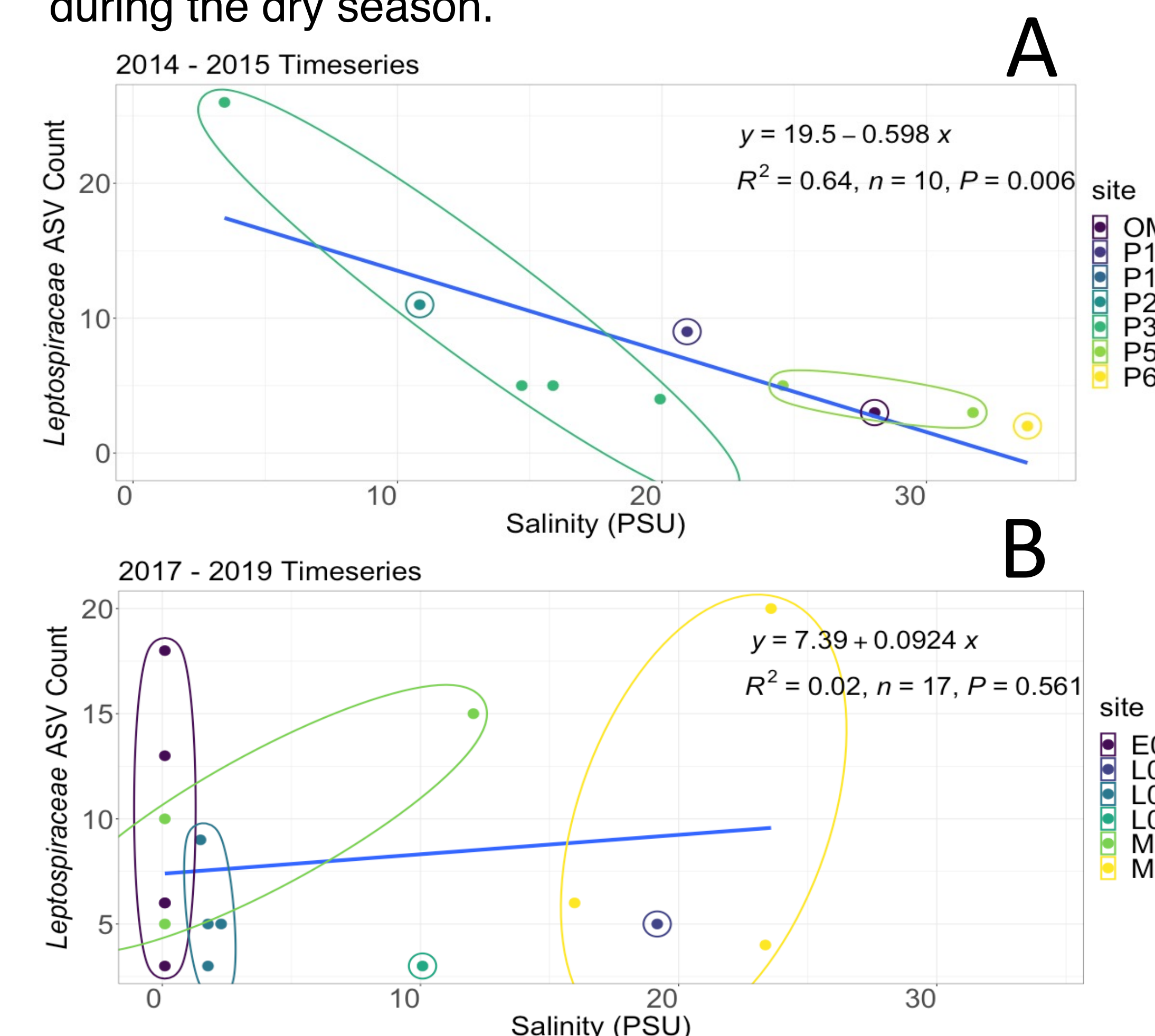


Figure 7: Correlation between salinity and the presence of *Leptospiraceae* in He'eia fishpond (2014-2015 : A; 2017-2019: B).