## Ziwei(Zoe) Wu

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#### **Education**

#### Sun Yat-Sen University (SYSU)

Aug. 2021 - Present

Master in Science

Major: Ecology (honors program: Evolutionary

Ecology)

Cumulative GPA: 3.5 (85.65/100)

Core Courses: Evolutionary Ecology, Seminar in Ecology

and Evolution, Data Analysis in R

#### Fujian Agriculture and Forestry University (FAFU)

Sep. 2017 - Jun. 2021

Bachelor of Agriculture, summa cum laude

Majors: Plant Science and Technology

Cumulative GPA: 3.4/4

Relevant Courses: Microbiology, Genetics (Bilingual

Course), Molecular Genetics, Biological Data Analysis, Organic Chemistry B, Plant Physiology A, Biochemistry B, Probability Theory, Experimental

Designs and Statistical Analyses, Plant

Cell Biology, Agroecology, Plant

Biotechnology

## Research Experience

1. Multi-omics study on the lignocellulose degradation by holobionts in mangrove sesarmid crabs: a game-

Nov. 2022 - Present

changer for terrestrialization

**Co-first author**, in progress, in collaboration with The Chinese University of Hong Kong

- Led the <u>assembly of three Sesarmidae genomes</u>.
- Conducted gene functional annotations, focusing on CAZymes, using public databases.

- Analyzed <u>genomic collinearity</u>, evaluated CAZymes families, and identified unique gene families and expansions/contractions.
- Combined <u>metagenomics</u> to uncover crab <u>lignocellulose degradation</u> mechanisms and evolution.

# 2. Evolution of terrestrialization in Land Crabs: exploring at the genomic level

Sep. 2022 - Present

First author, in progress, Master thesis

- Performed <u>chromosome-genome assembly</u> using ONT and Hi-C data.
- Extracted RNA from five tissues(Heart, Hepatopancreas, Testis, Muscles, Gills),
   self-constructed RNA libraries, and full-length cDNA ONT sequencing.
- Gene functional annotation.
- Analyzed genomic collinearity and identified unique gene families, expansions/contractions, and positive selection to understand its adaptive terrestrial evolution.

#### Orthomyxoviridae across metazoans

**Co-first author**, submitted to *PLOS Pathogens* in collaboration with The Hong Kong Polytechnic University

- Constructed the <u>phylogenetic tree</u> for the newly recognized viruses alongside known members of Orthomyxoviridae.
- Analyzed the ancestral states of the novel viruses and their host associations.
- 4. <u>Transcriptome Analysis of Pathogen-Induced</u>
  <u>Physiological Responses in Shrimp</u>

Feb. 2022 - Jan. 2023

First author, completed

- Gathered nine <u>RNA-Seq project datasets</u> from the NCBI SRA and finally obtained 109 transcriptome expression profiles.
- Data Quality Control, Differential Expression Analysis, and Enrichment Analysis (including Gene Ontology and Kyoto Encyclopedia of Genes Pathway Analysis)
- 5. <u>Microscopic Algae, "Macroscopic" Energy</u>

Jun. 2018 - Nov. 2021

## Student Leaders & Advisors, iGEM Project

- Constructed <u>transgenic algal strains</u> improve carbon dioxide fixation and lipid synthesis by overexpressing essential genes in the <u>ascorbic acid metabolic</u> pathway.
- Managed mathematical modeling efforts, including:
  - Creation of a Logistic growth model for microalgae based on growth trends.
  - Optimization models using cost and time data to predict optimal cultivation conditions, cost-effectiveness, and profitability of a microalgae facility.

#### **Publication** (\*Corresponding author; \*Co-first author)

- **1. Wu Z**, Chu KH\*, Ma KY\*. Transcriptome Analysis of Multiple Tissues in the Shrimp *Penaeus vannamei* Reveals the Typical Physiological Response to Three Pathogens[J]. Journal of Marine Science and Engineering. 2023; 11(2):389.
- 2. Veresoglou, S.D.\*, Chen, J<sup>+</sup>., **Wu, Z.**<sup>+</sup> et al. (2023). No-tillage outperforms conventional tillage under arid conditions and following fertilization. Soil Ecol. Lett. 5, 137–141 (2023).

#### Honors and Awards

- **1.** Gold and Bronze Awards at the International Genetically Engineered Machine (iGEM) in 2021 and 2019, respectively.
- **2.** Nominated for Best Sustainability (7/320 international student team) at iGEM in 2021
- 3. Second Prize Scholarship (8% students) in 2020
- 4. Advanced Individual in Social Work and Excellent Volunteer in Cangshan District

### Conference presentation

- **Wu ZW**, Chu KH, Ma KY. Transcriptome analysis of multiple tissues of *Penaeus vannamei* reveals the typical physiological response to the invasion of three pathogens, THE 16<sup>th</sup> SYMPOSIUM OF CRUSTACEAN SOCIETY, 12-13 Nov 2022, Heibei Province, China (oral presentation).
- **Wu ZW**, Chu KH, Ma KY. Transcriptome analysis of multiple tissues of *Penaeus vannamei* reveals the typical physiological response to the invasion of three pathogens, Dialogue with fantastic creatures—SYSU&HKU paleontology theme workshop, 15-17 Oct 2022, Guangdong Province, China (oral presentation).

## **Additional Experience**

1.	Undergraduate Ecological Comprehensive Experiment	Mar. 2023 - Jun. 2023
	Laboratory Assistant	
2.	Undergraduate Biochemical Experiment	Oct. 2022 - Jan. 2023
	Laboratory Assistant	
3.	English-Chinese translation of a sustainable development	Oct. 2022 - Nov. 2022
	report	
	United Nations Volunteers Translator	
4.	Undergraduate Ecological Comprehensive Experiment	Mar. 2022 - Jun. 2022
	Laboratory Assistant	

#### Skills and Hobbies

**Software:** Proficient in R (preferred for analysis and plotting), strong command of Perl programming (primary tools), familiar with Python (supplementary tools), limited exposure to Java

Hobbies: Avid Reader & Passionate Cook & Food Enthusiast & Animal Lover

## References

#### Ka Yan MA, Associate Professor

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#### Rongfeng Cui, Associate Professor

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#### **Tsang Ling Ming, Assistant Professor**

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The Chinese University of Hong Kong
<a href="mailto:lmtsang@cuhk.edu.hk">lmtsang@cuhk.edu.hk</a>

Stay curious forever, be willing to embrace everything, and excel at learning from scratch.