

Ziwei (Zoe) Wu

[Personal Homepage](#)

State Key Laboratory of Biocontrol, Southern Marine Science and Engineering Guangdong Laboratory (Zhuhai), School of Ecology, Sun Yat-sen University, Guangzhou, China

E-mail: ziweiw1998@gmail.com

Education

Sun Yat-sen University

Master in Science degree

Aug. 2021 - Present

Majors: **Ecology** (honors program: Evolutionary Ecology)

Cumulative GPA: 3.5 (85.65/100)

Anticipated Core Courses: Evolutionary Ecology, Seminar in Ecology and Evolution, Data Analysis in R

Fujian Agriculture and Forestry University

Bachelor of Agriculture degree, summa cum laude

Sep. 2017 - Jun. 2021

Majors: **Plant Science** and **Technology** (honors program, quantitative focus)

Cumulative GPA: 3.5 (85/100)

Relevant Courses: Organic Chemistry B, Microbiology, Plant Physiology A, Biochemistry B, Probability Theory, Biological Data, Experimental Designs and Statistical Analyses, Genetics (Bilingual Course), Molecular Genetics, Plant Cell Biology, Agroecology, Biological Data Analysis, Plant Biotechnology

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

Research Experience

1. Crabs and Symbiotic Microorganisms: Collaborative Cellulose Degradation

Facilitating Radiative Diversity in Sesarmidae — Genomic and Gut

Hongkong, China

Microbiome Macro-genomic Analysis of Crabs

Co-first author, In Analysis

Nov. 2022 - Present

- Led three Sesarmidae genome assemblies using third-generation ONT sequencing data.
- Conducted **gene functional annotation (especially the annotation of CAZymes)** using transcriptomic data and homologous protein information, utilizing the Maker tool and public database.
- ♦ Performed **three crabs' genomic collinearity analysis, CAZymes family assessment, and positive selection analysis** to identify distinctive gene families, significant expansions, and contractions in *Sesarmidae*.
- Combining metagenomics and metatranscriptomics to reveal the molecular mechanism and evolutionary process of crab degradation of lignocellulose.

2. Terrestrial Adaptation Evolution in Land Crabs: Exploring at the Genomic Level

Shenzhen, China

First author, Upcoming Graduation Thesis

Sep. 2022 - Present

- Completed **genome assembly** using third-generation ONT sequencing data and Hi-C data.
- Individually extracted RNA from five tissues of the Land Crabs (*Gecarcoidea lalandii*), **constructed RNA libraries**, and **performed independent third-generation full-length cDNA ONT sequencing**.
- Conducted **gene functional annotation** amalgamating second and third-generation transcriptomic data and homologous protein information.
- Performed inner **genomic collinearity analysis, unique and shared gene family assessment, and positive selection analysis** to identify distinctive gene families, significant expansions, and contractions in *Gecarcoidea lalandii*.

Additionally, scrutinized for genes under positive selection in *Gecarcoidea lalandii*, with the objective of elucidating the driving factors behind its adaptive evolution (terrestrial adaptation).

3. **Uncovering Influenza-Like Virus Clades and New Genera in Invertebrates: Evolutionary Insights into Orthomyxoviridae across Metazoans**

Hongkong, China

Co-first author, submitted in *Current Biology* (Cell Press)

Jun. 2022 - Jul. 2023

- Constructed **phylogenetic relationship** of newly identified viruses with known viral members from Orthomyxoviridae.
- Ancestral states** of the novel viruses and host associations analysis.

4. **Transcriptome Analysis of Pathogen-Induced Physiological Responses in Shrimp** Shenzhen, China

First author, Published

Feb. 2022 - Jan. 2023

- Gathered nine RNA-Seq project datasets from the NCBI SRA and finally obtained 109 transcriptome expression profiles.
- Generated **innovative ideas** to enhance project outcomes.
- Proficiently acquired expertise in **Shell scripting** and effectively utilized **R programming languages**, which enabled me to complete the Data Processing, Plot Generation, and Initial Draft Writing independently.

5. **Microscopic Algae, “Macroscopic” Energy**

Fuzhou, China

Student Leaders & Advisors, iGEM Project

Jun. 2018 - Nov. 2021

- Led the construction of transgenic algal strains to enhance carbon dioxide fixation and lipid synthesis metabolism by overexpressing critical genes in the ascorbic acid metabolic pathway.
- Led mathematical modeling experiments, **including**
 - Designing** a Logistic growth model for microalgae based on growth trends.
 - Developed** optimization models using cost and time data to predict optimal cultivation conditions and evaluate microalgae factory cost-effectiveness and profitability.

Academic Activities and Publication

- [1] **Wu ZW**, 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] **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 16th SYMPOSIUM OF CRUSTACEAN SOCIETY, 12-13 Nov 2022, Heibei Province, China (oral presentation).
- [3] **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 (Click for More Information)

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| 1. Undergraduate Ecological Comprehensive Experiment | Mar. 2023 - Jun. 2023 |
| Laboratory Assistant in the 2022 spring semester | High-Throughput Sequencing Module |
| 2. Undergraduate Biochemical Experiment | Oct. 2022 - Jan. 2023 |
| Laboratory Assistant in the 2022 autumn semester | Biochemical Experiment |
| 3. English - Chinese translation of a sustainable development report | Oct. 2022 - Nov. 2022 |
| United Nations Volunteers | Translator |
| 4. Undergraduate Ecological Comprehensive Experiment | Mar. 2022 - Jun. 2022 |
| Laboratory Assistant in the 2021 autumn semester | High-Throughput Sequencing Module |

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 & Cat Lover

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