

# YiFan Xiong

✉ [ewanxiong@gmail.com](mailto:ewanxiong@gmail.com)  [github.com/xyifan97](https://github.com/xyifan97)

## Education

**Fujian Agriculture and Forestry University** | *Fuzhou, Fujian Province, CN*

Sep. 2019 - Jun. 2022

- M.S. in Bioengineering. GPA: 3.33/4.0

**Wuhan Institute of Bioengineering** | *Wuhan, Hubei Province, CN*

Sep. 2015 - Jun. 2019

- B.S. in Bioengineering. GPA: 3.56/4.0

## Work experience

**Suzhou Dynamic Biosystems**

Jun. 2023 - Present

*Department of Bioinformatic, Bioinformatics Engineer*

- Independently developed **cellPCT**: An R package for single-cell percentage visualization
- **Well-Paired-Seq** platform single-cell data testing and troubleshooting.
- scRNA-seq and scVDJ-seq pipeline development and maintenance.

**Fuzhou Institute of Data Technology**

Jul. 2022 - Jun. 2023

*Department of Digital Medicine, Bioinformatics Engineer*

- Public data collection and normalization for hECA2.0 Single-cell database. (Collaborate with Tsinghua University)
- Multi-Omics pipeline construction and data analysis.

## Research Experience

**Undergraduate Research Assistant**

*Functional Bioactive Peptides Lab* | Supervisor: Prof. Junlin Zhang

Sep. 2016 - Jun. 2017

**Project 1: Extraction and identification of anti-cancer bioactive peptides from traditional Chinese medicine**

- Experiments including DNA extraction, PCR, vector construction, protein extraction, SDS-PAGE and cell culture, *et al.*

**Project 2: Functional verification of QinFeng tea by constructing Hyperuricemia animal models**

- Construct mouse Hyperuricemia models by two protocols: High-purine diet and injection of uric acid metabolism enzymes inhibitors.
- Models evaluate by blood sample collection and qualitative analysis of uric acid using ELISA.
- Data process and visualization.

*Plant Hormone and Nutrient Regulation Lab* | Supervisor: Prof. Zhongming Fang

Jul. 2017 - Jun. 2019

**Project 3: Altered Expression of *OsAAP3* Influences Rice Lesion Mimic and Leaf Senescence by Regulating Arginine Transport and Nitric Oxide Pathway (published)**

- *OsAAP3* transgenic rice leaf tissue RNA-seq data analysis, including QC, mapping, count calling, DEGs identification and functional enrichment analysis.
- Paper writing and data visualization.

**Graduate Student Researcher**

*Plant Functional genomics lab* | Supervisor: Prof. Shoukai Lin

Jun. 2019 - Jul. 2022

**Project 4: Genome-wide identification of the calcium-dependent protein kinase gene family in *Fragaria vesca* and expression analysis under different biotic stresses (published)**

- Determined the evolutionary history of FvCDPKs by genome-wide identification and collinearity analysis.
- Retrieved strawberry abiotic stress transcriptome data from public database.

- Different transcriptional patterns of wild strawberry under various abiotic stresses(Bacteria, Fungi, Virus).

**Project 5: Identification and Expression Analysis of CDPK Family in *Eriobotrya japonica*, reveals *EjCDPK25* in Response to Freezing Stress in Fruitlets(preprint)**

- Identification of CDPK gene family in loquat, basic gene family analysis including gene structure, protein motif, collinearity analysis.
- Transcription patterns of EjCDPKs and target gene identification by freezing stress traits associated co-expression genes.
- Target gene over-expression Arabidopsis germline cultivation and freezing stress tolerance experiments.

**Project 6: Large-scale Physiological and Transcriptome Analysis Insights into Adaptive Responses of *Eriobotrya japonica* Fruitlets to Freezing Stress(manuscript)**

- Genome-wide identification of Loquat protein kinase using plant's kinase HMM models and finding expression patterns of freezing-stress related protein kinase.

## Publications

---

- Wei Qilang, Zhenwei Yan, **Yifan Xiong**, and Zhongming Fang (2021). Altered Expression of OsAAP3 Influences Rice Lesion Mimic and Leaf Senescence by Regulating Arginine Transport and Nitric Oxide Pathway. *International Journal of Molecular Sciences*, 22, no.4: 2181. <https://doi.org/10.3390/ijms22042181>
- **Yifan Xiong**, Dahe Lin , Shiwei Ma, Chunhua Wang, Shoukai Lin (2022). Genome-wide identification of the calcium-dependent protein kinase gene family in fragaria vesca and expression analysis under different biotic stresses. *European Journal of Plant Pathology*. 164(2):283-98. <https://doi.org/10.1007/s10658-022-02560-4>
- **Yifan Xiong**, Shunquan Lin, Jincheng Wu, Shoukai Lin (2024). Identification and Expression Analysis of CDPK Family in *Eriobotrya japonica*, reveals *EjCDPK25* in Response to Freezing Stress in Fruitlets. *bioRxiv*. <https://doi.org/10.1101/2024.05.01.591999>

## Research Skills

---

### Bioinformatic skills

- Programming languages: R, Python and Shell
- NGS data analysis: WGS, WES, scRNA-seq, bulk RNA-seq(miRNA, lncRNA, circRNA), WGBS, ChIP-seq, ATAC-seq
- Workflow Management: WDL, Snakemake
- Code Management and Version Control: Github and Gitlab
- Servers and Environment Manager: Conda, Docker, Slurm

### Other skills

- Vector construction, Real-time PCR, SDS-PAGE, Cell line cultivation, plant over-expression material construction
- Thesis writing and data visualization
- Oral presentation and poster presentations
- Independent thinking ,problem-solving, Team working and communication
- English proficiency: IETLS 7.0 (Listening 7.5, Reading 8.5, Writing 6.0, Speaking 6.5)

## Academic activities

---

<b>The 10th National Symposium on Loquat</b>   <i>Lanxi, Zhejiang province</i>	<b>May. 2021</b>
<ul style="list-style-type: none"><li>• Submit an abstract on the topic of <i>Identification calcium sensors in loquat</i></li></ul>	
<b>Academic poster competition for Graduate Student</b>   <i>Fujian Agriculture and Forestry University</i>	<b>May. 2022</b>
<ul style="list-style-type: none"><li>• Oral and poster presentation on the topic of my graduate thesis</li></ul>	
<b>Digital Medicine Algorithm Innovation Competition</b>   <i>Fuzhou, Fujian province</i>	<b>Aug. 2022</b>
<ul style="list-style-type: none"><li>• Organization assistant</li></ul>	
<b>The 2nd Symposium of Fujian Bioinformation Society</b>   <i>Fuzhou, Fujian province</i>	<b>Feb. 2023</b>
<ul style="list-style-type: none"><li>• Poster presentation</li></ul>	

## Honors & Awards

---

<b>Scholarship for Undergraduates 3rd Place</b>   <i>Wuhan Institute of Bioengineering</i>	<b>Dec. 2015</b>
<b>Annual Outstanding Undergraduates</b>   <i>Wuhan Institute of Bioengineering</i>	<b>Dec. 2016</b>
<b>National English Competition for Undergraduates 3rd Place</b>   <i>Wuhan Institute of Bioengineering</i>	<b>May. 2017</b>
<b>Scholarship for Undergraduates 2nd Place</b>   <i>Wuhan Institute of Bioengineering</i>	<b>Dec. 2017</b>
<b>Scholarship for Undergraduates 1st Place</b>   <i>Wuhan Institute of Bioengineering</i>	<b>Dec. 2018</b>
<b>Outstanding Graduates</b>   <i>Wuhan Institute of Bioengineering</i>	<b>Jun. 2019</b>
<b>Scholarship for Graduates</b>   <i>Fujian Agriculture and Forestry University</i>	<b>Jun. 2020</b>
<b>Academic Achievements Prize for Graduates</b>   <i>Fujian Agriculture and Forestry University</i>	<b>Jun. 2022</b>