YiFan Xiong

Education

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

Sep. 2019 - Jun. 2022

- M.S. in Bioengineering. GPA: 3.33/4.0
- Focus area on Functional genomics of plant.

Wuhan Institute of Bioengineering | Wuhan, Hubei Province, CN

Sep. 2015 - Jun. 2019

- B.S. in Bioengineering. GPA: 3.56/4.0
- Lab rotation experience in experimental and bioinformatic.

Work experience

Suzhou Dynamic Biosystems

Jun. 2023 - Present

Department of Bioinformatic, Bioinformatics Engineer

• Independently developed cellPCT: An R package for single-cell percentage visulization

Fuzhou Institue of Data Technology

Jul. 2022 - Jun. 2023

Department of Digital Medicine, Bioinformatics Engineer

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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 gutitive analsis of uric acid using ELISA.
- Data process and visulization.

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 writting and data visulization.

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 strawbarry abiotic stress transciptome 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, Chlp-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 writting and data visulization
- Oral presentaion and poster presentations
- Independent thinking ,problem-solving, Team working and communication
- English competency: IETLS 7.0 (Listening 7.5, Reading 8.5, Writing 6.0, Speaking 6.5)

Acdemic activates

The 10th National Symposium on Loquat Lanxi, Zhejiang province	May. 2021
Submit an abstract	
Academic poster competition for Graduate Student Fujian Agriculture and Forestry University	May. 2022
Oral and poster presentation on the topic of my gradute thesis	
Digital Medicine Algorithm Innovation Competition Fuzhou, Fujian province	Aug. 2022
Organization assistant	
The Second Symposium of Fujian Bioinformation Society Fuzhou, Fujian province	Feb. 2023
Poster presentation	
Honors & Awards	
Scholarship for Undergraduates 3rd Place Wuhan Institute of Bioengineering	Dec. 2015
Annual Outstanding Undergraduates Wuhan Institute of Bioengineering	Dec. 2016
National Undergraduates' English Competition 3rd Place Wuhan Institute of Bioengineering	May. 2017
Scholarship for Undergraduates 2nd Place Wuhan Institute of Bioengineering	Dec. 2017
Scholarship for Undergraduates 1st Place Wuhan Institute of Bioengineering	Dec. 2018
Scholarship for orderigraduates 1st Flace Warran Institute of Dioengineering	
Outstanding Graduates Wuhan Institute of Bioengineering	Jun. 2019
	Jun. 2019 Jun. 2020