Yifan Xiong

Education

Fujian Agriculture and Forestry University | M.S. in Bioengineering.

Sep. 2019 - Jun. 2022

Wuhan University of Bioengineering | B.S. in Bioengineering.

Sep. 2015 - Jun. 2019

Professional experience

Bioinformatics engineer | Dynamic Biosystems Ltd, Xiamen, China

Jun. 2023 - Jun. 2024

Development team member of the Well-Paired-Seq single-cell platform; Independently developed R package cellPCT.

Bioinformatics engineer | Tsinghua-Fuzhou Institue of Data Technologies, Fuzhou, China

Jul. 2022 - Jun. 2023

• Metadata organization and cell type annotation for the hECA2.0 Single-cell Atlas; Construct Multi-omics pipelines.

Research Experience

Bioinformatics Research Assistant

Jul. 2024 - Present

Department of Automation, Tsinghua University, PI: Prof. Xuegong Zhang

Evaluate the biological meaning in different single cell foundation models and LLM's gene embedding

• Extract gene embedding from different models, evaluate the biological meaning by multi downstream tasks.

Reversing T cell exhaustion through TF combinations identified by scDirect algorithm

• Annotate T cell subclusters in scRNA-seq datasets and deduce TF combinations for reversing exhausted T cells.

LncRNA and ceRNA analysis of Marfan syndrome reveal the potential role of lncRNA in the pathogenesis

Co-expression analysis of lncRNA and mRNA identifies potential pathogenesis ncRNAs involved in MF syndrome.

Graduate Student Researcher

Jun. 2019 - Jul. 2022

Department of Life Science, Fujian Agriculture and Forestry University, PI: Prof. Shoukai Lin

Identify Calcium-dependent protein kinase in *Fragaria vesca* and reveal their expression patterns under biotic stresses

- Phylogenetic analysis of CDPK family in strawberry, and determaine the stress-related Cis-regulatory elements.
- Conduct transcriptome analysis of different pathogen infections to reveal the distinctive functional responses.

Large-scale physiological and transcriptome analysis of adaptive responses of *Eriobotrya japonica* under freezing stress

- · Genome-wide identification of Loquat protein kinase and TFs using motif HMM models and sequence similarity.
- Associate the frezzing-stress traits with co-expressed gene modules, find the hub genes in gene regulation networks.
- Experimental verification of target genes in Arabidopsis, revealing the function of freezing stress resistance.

Undergraduate Research Assistant

Sep. 2016 - Jun. 2019

Applied Biotechnology Research Center, Wuhan University of Bioengineering, Pl: Prof. Junlin Zhang & Prof. Zhongming Fang

Extraction and identification of anti-cancer bioactive peptides and in vitro functional verification

• Wet-lab training including vector construction, QPCR, SDS-PAGE, cell culture, annimal model establishment.

Transcriptome analysis of *Oryza sativa* transgenic materials reveal mechanism of leaf senescence

• OsAAP3 transgenic rice leaf tissue RNA-seq data analysis, conduct data visualization, and respond to peer reviews.

RNA-seg analysis provides insights into the regulatory mechanism of striped leaf albinism in *Areca catechu*

• *De novo* transcriptome analysis of arecanut leaf, determaine the expression patterns of genes related to pigments biosynthesis, senescence and chloroplast development.

Publications

- Yifan Xiong, Shunquan Lin, Jincheng Wu, Shoukai Lin. Identification and Expression Analysis of CDPK Family in Eriobotrya japonica, reveals EjCDPK25 in Response to Freezing Stress in Fruitlets. bioRxiv, 2024. https://doi.org/10.1101/2024.05.01.591999
- Yifan Xiong, Dahe Lin, Shiwei Ma, Chunhua Wang, Shoukai Lin. 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, 2022, 164(2):283-98. https://doi.org/10.1007/s10658-022-02560-4
- Wei Qilang, Zhenwei Yan, **Yifan Xiong**, and Zhongming Fang. Altered Expression of *OsAAP3* Influences Rice Lesion Mimic and Leaf Senescence by Regulating Arginine Transport and Nitric Oxide Pathway. *International Journal of* Molecular Sciences, 2021, 22, no.4: 2181. https://doi.org/10.3390/ijms22042181
- Jia Li, Xiaocheng Jia, Liyun Liu, Xianmei Cao, **Yifan Xiong**, Yaodong Yang, Huangi Zhou, Ming Yi, and Meng Li. "Comparative biochemical and transcriptome analysis provides insights into the regulatory mechanism of striped leaf albinism in arecanut (Areca catechu L.)." Industrial Crops and Products, 2020, 154: 112734. https://doi.org/10.1016/j.indcrop.2020.112734

Skills

- Language: IELTS overall band 7 (Reading 8.5, Listening 7.5, speaking 6.5, writing 6.0), native in mandrain Chinese.
- Programming: Proficient in R, Python and Shell
- NGS data analysis: scRNA-seq, bulk RNA-seq(ceRNA competition), WGS, WGBS, ATAC-seq
- Code Management and Version Control: Github and Gitlab
- Servers, Environment and Workflow Manager: Slurm, Conda, Docker, WDL and Snakemake

Presentations and Posters

The 2nd Symposium of Fujian Bioinformation Society Poster presentation Fuzhou, Fujian province	Feb. 2023
Digital Medicine Algorithm Innovation Competition Organization assistant Fuzhou, Fujian province	Aug. 2022
Academic competition for Graduate Student Oral and poster presentation Fuzhou, Fujian province	May. 2022
The 10th National Symposium on Loquat Poster presentation Lanxi, Zhejiang province	May. 2021
Honors & Awards	
Employee of the Year Award Dynamic Biosystems Ltd.	Dec. 2023
Academic Achievements Prize for Graduates Fujian Agriculture and Forestry University	Jun. 2022
Scholarship for Graduates Fujian Agriculture and Forestry University	Jun. 2020
Outstanding Graduates Wuhan University of Bioengineering	Jun. 2019
Scholarship for Undergraduates 1st Place Wuhan University of Bioengineering	Dec. 2018
Scholarship for Undergraduates 2nd Place Wuhan University of Bioengineering	Dec. 2017
National English Competition for Undergraduates 3rd Place Wuhan University of Bioengineering	May. 2017
Annual Outstanding Undergraduates Wuhan University of Bioengineering	Dec. 2016
Scholarship for Undergraduates 3rd Place Wuhan University of Bioengineering	Dec. 2015

Outreach and Extracurricular Activities

Community volunteer for anti-COVID pandemic	Jun. 2020
Bilingual volunteer for the Tour of China Cycling Competition	May. 2016
Pre-College Summer School Program @Central Chnia Normal University	Jul. 2015 - Aug. 2015