

# Liaoyi (Luis) Xu

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## EDUCATION

- 2022 – Present    **Ph.D. in Bioinformatics and Computational Biology**  
Department of Integrative Biology  
University of Texas at Austin, TX, USA  
Advisor: [Arbel Harpak](#)
- 2018 – 2021    **M.A. in Animal Science**  
Zhejiang University, Hangzhou, China
- 2014 – 2018    **B.A. in Life Science and Technology**  
Southwest University for Nationalities, Chengdu, China

## HONORS & AWARDS

- 2023 – 2024    UT Austin Professional Development Awards
- 2023    ASHG 2023 Charles Epstein Award Finalist (**Top 2%**)
- 2023    American Chinese Geneticist Association (ACGA) Trainee Award
- 2021    Excellent Postgraduate Students' Award (merit honor for **Top 10%**)
- 2020    Sunshine Scholarship (**Top 3** students in the major)
- 2020    Graduate of Merit/Triple A Graduate (merit honor for **Top 5%**)
- 2020    Award of Honor for Graduate (merit honor for **Top 10%**)
- 2018 – 2020    Academic Scholarship
- 2018    Excellent Graduate Students' Award (merit honor for **Top 10%**)
- 2014 – 2018    The Agriculture & Forestry Scholarship

## PUBLICATIONS

- 2025    Liaoyi Xu, Eucharist Kun, Devansh Pandey, Joyce Y Wang, Marianne F Brasil, Tarjinder Singh, and Vagheesh M Narasimhan. “**The genetic architecture of and evolutionary constraints on the human pelvic form**”. *Science*
- 2023    Yanhong Chen, Wei Cheng, Jia Tang, Boneng Xiao, Ming Kuang, Liaoyi Xu, and Hongying Liu. “**Mitigating cognitive impairment in aging mice: Exploring the therapeutic potential of ischelim**”. *Biochemical and Biophysical Research Communications*
- 2022    Tingyao Zhu, Liaoyi Xu, Jiacheng Peng, Ming Chen, and Haisheng Xu. “**Molecular characteristics and immune function of ubiquitin C-terminal hydrolase-L3 in *Macrobrachium nipponense***”. *Fish & Shellfish Immunology*

- 2021 Liaoyi Xu, Xiefei Zhou, Peichen Wang, Qian Jin, Tingyao Zhu, Ming Chen, and Haisheng Xu. **“The novel six LIM and one PET domain-containing protein Lmpt is involved in the immune response through activation of the NF- $\kappa$ B signalling pathway in the crustacean, *Macrobrachium nipponense*”**. Developmental & Comparative Immunology
- 2021 Wei Zhang, Qiangqiang Chen, Liaoyi Xu, Jianfeng Cai, and Jinzhi Zhang. **“The potential role of PSMA6 in modulating fat deposition in pigs by promoting preadipocyte proliferation and differentiation”**. Gene
- 2021 Tingyao Zhu, Xiefei Zhou, Wei Zhang, Yue Wu, Jingjing Yang, Liaoyi Xu, Ming Chen, Weiren Dong, and Haisheng Xu. **“Multiplex and real-time PCR for qualitative and quantitative donkey meat adulteration”**. Journal of Food Measurement and Characterization
- 2020 Liaoyi Xu, Yue Wu, Xiefei Zhou, Peichen Wang, and Haisheng Xu. **“Characterization and immune function of decapentaplegic (Dpp) gene from the oriental river prawn, *Macrobrachium nipponense*”**. Fish & Shellfish Immunology
- 2020 Liaoyi Xu, Xiefei Zhou, Yue Wu, Jingjing Yang, and Haisheng Xu. **“A novel SNW/SKIP domain-containing protein, Bx42, is involved in the antibacterial responses of *Macrobrachium nipponense*”**. Developmental & Comparative Immunology
- 2020 Yue Wu, Yuxiao Sun, Wanrong Zhang, Jingjing Yang, Xiefei Zhou, Liaoyi Xu, Peichen Wang, and Haisheng Xu. **“Molecular cloning, characterization, and expression analysis of lipopolysaccharide-induced TNF- $\alpha$  factor (LITAF) in the mud crab, *Scylla paramamosain*”**. Crustaceana

## CONFERENCE PRESENTATIONS

- 2023 Liaoyi Xu, Eucharist Kun, Marianne F. Brasil, Tarjinder Singh, Vagheesh M. Narasimhan. **Deep learning to understand the genetic architecture and evolution of the human pelvis**. *Society for Molecular Biology and Evolution (SMBE)*, Puerto Vallarta, Mexico (Poster Presentation)
- 2023 Liaoyi Xu, Eucharist Kun, Marianne F. Brasil, Tarjinder Singh, Vagheesh M. Narasimhan. **Deep learning to understand the genetic architecture and evolution of the human pelvis**. *American Society of Human Genetics (ASHG) Annual Meeting*, Washington, D.C., USA (Oral Presentation)

## TALKS

- 2024 **Deep learning to understand the genetic architecture and evolution of the human pelvis**. *ACGA Virtual Seminar* (Invited)
- 2023 **Deep learning to understand the genetic architecture and evolution of the human pelvis**. The University of Texas at Austin, *Special BASIL Meeting*, Department of Anthropology
- 2023 **Deep learning to understand the genetic architecture and evolution of the human pelvis**. The University of Texas at Austin, *BioTACOs*, Department of Integrative Biology

## SKILLS

Computational: Python (Pandas, Polars, Matplotlib, Seaborn, PyTorch, pyMC, Numpy, Scikit-learn, JAX, Statsmodels), R (JAGS, Tidyverse, RStan, ggplot2), SQL,  $\text{\LaTeX}$ , UNIX shell.

Experimental: RNA/DNA extraction and purification, PCR/qPCR, molecular cloning, siRNA and dsRNA interference, CRISPR related gene-editing, cell culture, flow cytometry, Western blot.

Languages: Mandarin (Native), English (Proficient).