Qisheng Pan

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RESEARCH INTEREST

I studied human genetic variants, investigating the relationship between protein mutations and pathogenic phenotypes. By leveraging computational biophysical measurements and machine learning, my research focuses on providing a guideline of the interpretation of current methods, and the development of new approaches to better characterise the effect of variants.

EDUCATION

University of Queensland

• Doctor of Philosophy (Computational Biology)

University of Melbourne

• Master of Science (Bioinformatic)

Brisbane, Australia

Jun 2025 (expected)

Melbourne, Australia

2019-2021

South China Normal University

• Bachelor of Science (Biotechnology)

Guangzhou, China *2015-2019*

PUBLICATIONS

- 1. **Pan Q**, Parra G, Myung Y et al. AlzDiscovery: A computational tool to identify Alzheimer's disease-causing missense mutations using protein structure information, Protein Sci 2024.
- 2. **Pan Q**, Portelli S, Nguyen TB et al. Characterization on the oncogenic effect of the missense mutations of p53 via machine learning, Brief Bioinform 2023;25.
- 3. Serghini A, Portelli S, Troadec G et al. Characterizing and predicting ccRCC-causing missense mutations in Von Hippel-Lindau disease, Hum Mol Genet 2023.
- 4. Jessen-Howard D, **Pan Q**, Ascher DB. Identifying the Molecular Drivers of Pathogenic Aldehyde Dehydrogenase Missense Mutations in Cancer and Non-Cancer Diseases, Int J Mol Sci 2023;24.
- 5. Zhou Y, **Pan Q**, Pires DEV et al. DDMut: predicting effects of mutations on protein stability using deep learning, Nucleic Acids Res 2023;51:W122-W128.
- 6. Boer JC, **Pan Q**, Holien JK et al. A bias of Asparagine to Lysine mutations in SARS-CoV-2 outside the receptor binding domain affects protein flexibility, Front Immunol 2022;13:954435.
- 7. **Pan Q**, Nguyen TB, Ascher DB et al. Systematic evaluation of computational tools to predict the effects of mutations on protein stability in the absence of experimental structures, Brief Bioinform 2022;23.
- 8. Han YY, Jin K, **Pan QS** et al. Microglial activation in the dorsal striatum participates in anxiety-like behavior in Cyld knockout mice, Brain Behav Immun 2020;89:326-338.

TEACHING EXPERIENCES

Teaching Assistant, University of Queensland

Aug 2024

• BIOT7060: Frontiers in Medical Biotechnology

Instructors, University of Queensland

Nov 2023

• Advanced Data Visualisation with *ggplot2*: This workshop is the one that I designed, prepared, and delivered, focusing on practical skills on presenting data using *R* and *ggplot2* package.

Teaching Assistant, University of Queensland

Sept 2022

• Computing4lifescience Series

MENTORSHIP

Research Supervisor (UG: undergraduate, MS: master's)

Georgina Becerra Parra (UG, 2022, UQ), Dana Jessen-Howard (MS, 2023, UQ), Joshua Khoo (MS, 2024, UQ)

HONOURS & AWARDS

Conference support of SAAFE 2024 AMR Solutions Summit	
• Travel Awards of MM2023 conference	Dec 2023
• SCMB Award for Outstanding Contribution to Research (Group Awards)	Nov2023
• Student Prize in the CTCMS Seminar	Mar 2023
Comprehensive Student Scholarship	Sept 2016

COLLABORATIONS

Jeniffer Boer and Magdalena Plebanski, Royal Melbourne Institute of Technology, Australia

2022 - 2023

• Investigating the variants of the Spike protein in Omicron SARS-CoV-2 virus

RESEARCH EXPERIENCES

Characterising the pathogenic effect of missense mutations via machine learning

2022 - now

- · Leveraged different computational biophysical measurements to annotate missense variants.
- Developed machine learning models to classify phenotypes of mutations.

Benchmarking computational biophysical measurements in the absence of experimental 2022 - now structures

- Built high-throughput pipeline to generate protein homology models and AlphaFold models.
- Used different metric to assess the predictive performance of various machine learning models.

PRESENTATIONS

1. Poster presentation in the SAAFE 2024 AMR Solutions Summit (Australia)	Sept 2024
2. Poster presentation in the Lorne Protein Conference 2024 (Australia)	Feb 2024
3. Oral and poster presentations in the MM2023 conference (Australia)	Dec 2023
4. Poster presentation in the ABACBS 2023 conference (Australia)	Dec 2023
5. Research Talk in the 22 nd International Conference on Bioinformatics (Australia)	Nov 2023
6. Lighting talk in the GenGen seminar (UQ)	Apr 2023
7. Oral presentations in the CTCMS seminar (UQ)	
8. Poster presentations in the Lorne Protein Conference 2023 (Australia)	
9. Poster presentation in the 18 th Annual Research Student Symposium (UQ)	Nov 2022
10. Oral presentations in the Joint Biomolecular and Medicinal Chemistry Theme Symposium (UQ)	Apr 2022

TECHNICAL SKILLS

Programming: Python, R, Linux Bash, JavaScript

Software: BLAST, MODELLER, PyMol, AutoDock Vina, GALAXY, etc. Machine learning: Random Forest, Neural Network, Feature selection, etc.

REFEREES

David B. Ascher	Professor, University of Queensland	d.ascher@uq.edu.au
Thanh-Binh Nguyen	Research Fellow, University of Queensland	thanhbinh.nguyen@uq.edu.au
Stephanie Portelli	Research Fellow, University of Queensland	s.portelli@uq.edu.au
Douglas E.V. Pires	Senior lecturer, University of Melbourne	douglas.pires@unimelb.edu.au
Cheng Long	Professor, South China Normal University	longcheng@m.scnu.edu.cn