

# Qisheng Pan

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## RESEARCH INTEREST

I applied knowledge-based modelling to study complex biological and chemical questions. In particular, I model enzyme-reaction interaction using machine learning approaches, which helps characterise enzyme function and enhance enzyme engineering.

## EDUCATION

|  |                      |
|--|----------------------|
| <b>University of Queensland</b>                | Brisbane, Australia  |
| • Doctor of Philosophy (Computational Biology) | 2022-2025            |
| <b>University of Melbourne</b>                 | Melbourne, Australia |
| • Master of Science (Bioinformatics)           | 2019-2021            |
| <b>South China Normal University</b>           | Guangzhou, China     |
| • Bachelor of Science (Biotechnology)          | 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

|   |           |
|---|-----------|
| <b>Tutor</b> , University of Queensland   | Feb 2025  |
| • <b>BINF6000: Bioinformatics Introduction</b>  |           |
| <b>Teaching Assistant</b> , University of Queensland  | Aug 2024  |
| • <b>BIOT7060: Frontiers in Medical Biotechnology</b>   |           |
| <b>Instructor</b> , University of Queensland  | Nov 2023  |
| • <b>Advanced Data Visualisation with ggplot2</b> : This workshop is the one that I designed, prepared, and delivered, focusing on practical skills on presenting data using R and ggplot2 package. |           |
| <b>Teaching Assistant</b> , University of Queensland  | Sept 2022 |
| • <b>Computing4lifescience Series</b>   |           |

## 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), Wuyang Ren (MS, 2025, UQ), Mingze Xu (MS, 2025, UQ)

## HONOURS & AWARDS

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|---|------------------|
| • Selected Yong Scientist Participants of Hong Kong Laureate Forum 2025 | <i>Nov 2025</i>  |
| • Conference support of SAAFE 2024 AMR Solutions Summit (\$1000)        | <i>Sept 2024</i> |
| • Travel Awards of MM2023 conference (\$300)                            | <i>Dec 2023</i>  |
| • SCMB Award for Outstanding Contribution to Research (Group Awards)    | <i>Nov 2023</i>  |
| • Student Prize in the CTCMS Seminar                                    | <i>Mar 2023</i>  |
| • Comprehensive Student Scholarship (\$500)                             | <i>Sept 2016</i> |

## COLLABORATIONS

|  |                    |
|--|--------------------|
| <b>Jeniffer Boer and Magdalena Plebanski, Royal Melbourne Institute of Technology, Australia</b> | <i>2022 - 2023</i> |
| • Investigating the variants of the Spike protein in Omicron SARS-CoV-2 virus                    |                    |

## RESEARCH EXPERIENCES

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|--|-------------------|
| <b>Characterising the pathogenic effect of missense mutations via machine learning</b>               | <i>2022 - now</i> |
| • Leveraged different computational biophysical measurements to annotate missense variants.          |                   |
| • Developed machine learning models to classify phenotypes of mutations.                             |                   |
| <b>Benchmarking computational biophysical measurements in the absence of experimental structures</b> | <i>2022 - now</i> |
| • 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

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|---|------------------|
| 1. Poster presentation in the ISMB 2025 Conference (United Kingdom)                             | <i>July 2024</i> |
| 2. Poster presentation in the Lorne Protein 2025 Conference (Australia)                         | <i>Nov 2024</i>  |
| 3. Poster presentation in the ABACBS 2024 Conference (Australia)                                | <i>Nov 2024</i>  |
| 4. Poster presentation in the SAAFE 2024 AMR Solutions Summit (Australia)                       | <i>Sept 2024</i> |
| 5. Poster presentation in the Lorne Protein 2024 Conference (Australia)                         | <i>Feb 2024</i>  |
| 6. Oral and poster presentations in the MM2023 Conference (Australia)                           | <i>Dec 2023</i>  |
| 7. Poster presentation in the ABACBS 2023 Conference (Australia)                                | <i>Dec 2023</i>  |
| 8. Research Talk in the 22 <sup>nd</sup> International Conference on Bioinformatics (Australia) | <i>Nov 2023</i>  |
| 9. Lighting talk in the GenGen seminar (UQ)   | <i>Apr 2023</i>  |
| 10. Oral presentations in the CTCMS seminar (UQ)  | <i>Mar 2023</i>  |
| 11. Poster presentations in the Lorne Protein 2023 Conference (Australia)                       | <i>Feb 2023</i>  |
| 12. Poster presentation in the 18 <sup>th</sup> Annual Research Student Symposium (UQ)          | <i>Nov 2022</i>  |
| 13. Oral presentations in the Joint Biomolecular and Medicinal Chemistry Theme Symposium (UQ)   | <i>Apr 2022</i>  |

## TECHNICAL SKILLS

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Programming: Python, R, Linux Bash, JavaScript

Software: BLAST, MODELLER, PyMol, AutoDock Vina, GALAXY, etc.

Machine learning: Random Forest, Neural Network, Feature selection, Transformer etc.

## REFEREES

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