

Hui Jia Farm

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RESEARCH INTEREST

Cardiac electrophysiology, Computational biology

ACADEMIC QUALIFICATIONS

Doctor of Philosophy, Computer Science

Oct 2020 –

University of Oxford, United Kingdom

Title: Computational Modelling of Ion Channel Kinetics under Drug Intervention on Cardiac Myocytes

Supervisors: Prof. David J. Gavaghan, Dr. Michael Clerx, Dr. Fergus Cooper, Dr. Chon Lok Lei, Dr. Ken Wang, Dr. Liudmila Polonchuk

Bachelor of Science (Mathematical Sciences), Honours

Aug 2014 – Jun 2018

Nanyang Technological University, Singapore

Specialisation: Applied Mathematics

Thesis: Stochastic Gradient Algorithm for Gaussian Graphical Models with Hidden Variables

Supervisors: Assoc. Prof. Xiang Liming, Assoc. Prof. Justin Dauwels

HONOURS AND AWARDS

A*STAR National Science Scholarship (PhD)

2018 –

ASEAN Undergraduate Scholarship

2014 – 2018

RESEARCH EXPERIENCE

Research Officer

Sep 2018 – Aug 2020

*Bioinformatics Institute, Agency for Science Technology and Research (A*STAR), Singapore*

- Implemented image processing techniques and developed various mathematical models to study biological phenomena, such as cell stretching and merozoite cell division.

Research Assistant

Jul 2018 – Aug 2018

Earth Observatory of Singapore, Nanyang Technological University, Singapore

- Solved inverse problems, with data uncertainties, non-uniqueness and theoretical uncertainties, to determine source of earthquakes from several signal stations.

Final Year Project Student

Aug 2017 – May 2018

School of Physical and Mathematical Sciences, Nanyang Technological University, Singapore

- Developed stochastic gradient algorithm to learn the structure of Gaussian graphical models with hidden variables at lower computational complexity by implementing sampling and projection method for estimation of gradient.

Research Intern

May 2017 – Jul 2017

*Bioinformatics Institute, Agency for Science Technology and Research (A*STAR), Singapore*

- Improved accuracy of Traction Force Microscopy algorithm, for cell biophysics, by 20% for low noise level data, with L1 and L2 regularization.

TEACHING EXPERIENCE

Marker

Oct 2021 – Dec 2021

Department of Computer Science, University of Oxford, United Kingdom

Course: Bayesian Statistical Probabilistic Programming

Demonstrator

Oct 2021 – Oct 2021

Doctoral Training Centre, University of Oxford, United Kingdom

Course: Software Engineering & Sustainable Research

Tutor

Aug 2017 – Nov 2017

School of Physical and Mathematical Sciences, Nanyang Technological University, Singapore

Course: Linear Algebra 1

OTHER ACTIVITIES

Bioinference 2022 Conference Organising Committee

Oct 2021 – May 2022

Funded by the London Mathematical Society, the Heilbronn Institute for Mathematical Research and the Department of Computer Science, University of Oxford

<https://bioinference.github.io/2022/>

Squash Club

May 2021 –

Captain of Green Templeton Squash Club, University of Oxford (2022 – 2023)

President of Green Templeton Squash Club, University of Oxford (2023 – present)

PUBLICATIONS

Farm, H.J., Clerx, M., Cooper, F. et al. (2023) “Importance of modelling hERG binding mechanism in predicting drug-induced action potential prolongations for drug safety assessment.” *Front. Pharmacol.* 14.

[doi:10.3389/fphar.2023/1110555](https://doi.org/10.3389/fphar.2023/1110555)

<https://github.com/FarmHJ/importance-of-binding-mechanism>

Creswell, R.*, Augustin, D.*, Bouros, I.*, **Farm, H.J.***, Miao, S.*, Ahern, A.* et al. (2022) “Heterogeneity in the onwads transmission risk between local and imported cases affects practical estimates of the time-dependent reproduction number.” *Phil. Trans. R. Soc. A.* 380:20210308. (*shared first authors) [doi:10.1098/rsta.2021.0308](https://doi.org/10.1098/rsta.2021.0308)

<https://github.com/SABS-R3-Epidemiology/transmission-heterogeneity-results>

van der Vegt, S. A.*, Dai, L.*, Bouros, I.*, **Farm, H.J.***, Creswell, R.*, Dimdore-Miles, O.*, Cazimoglu, I.* et al. (2022) “Learning transmission dynamics modelling of COVID-19 using comomodels.” *Math. Biosci.* 349.

(*shared first authors) [doi:10.1016/j.mbs.2022.108824](https://doi.org/10.1016/j.mbs.2022.108824)

<https://github.com/Como-DTC-Collaboration/como-models-math-biosci>

Zhou, T., ..., **Farm, H.J.**, Goh, E. L. K., Chiam, K. H. (2023) “ContrastivePose: A contrastive learning approach for self-supervised feature engineering for pose estimation and behavioral classification of interacting animals.” *Comput. Biol. Med.* 165:107416. [doi:10.1016/j.combiomed.2023.107416](https://doi.org/10.1016/j.combiomed.2023.107416)

Chong, L. H., Ching, T., **Farm, H.J.**, et al. (2022) “Integration of a microfluidic multicellular coculture array with machine learning analysis to predict adverse cutaneous drug reactions.” *Lab Chip.* 22, 1890-1904.

[doi:10.1039/D1LC01140E](https://doi.org/10.1039/D1LC01140E)

CONFERENCES

Potential importance of modelling hERG kinetics in predicting drug-induced action potential and QTc change: exploration through a simulation study informed by experimental data

12 Sep 2022

Rapid Fire Poster Presentation at Safety Pharmacology Society (SPS) Annual Meeting 2022, Montreal, Canada

Predicted drug effects on the action potential vary significantly with different models

14 Feb 2022

Poster Presentation at Biophysical Society (BPS) Annual Meeting 2024, Philadelphia, Pennsylvania, U.S.

Last Updated on 26 Jan 2024