

# Lee Jin Wee

✉ jinweelee@gmail.com | 🌐 thecausalclinician.com | in lee-jin-wee-425b1b13a | 📄 Google Scholar

## Education and Achievements

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### National University of Singapore

BACHELOR OF SCIENCE (HONOURS) IN COMPUTATIONAL BIOLOGY

2015-2019

- **Honour Roll**, University Scholars Programme — *Academic Year 2015/2016*
- **Deans List**, Faculty of Science — *Academic Year 2016/2017*

MASTERS OF SCIENCE IN STATISTICS

2020-2021

### Duke-NUS Medical School

DOCTOR OF MEDICINE

2023-2027

## Undergraduate Research Experience

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### Undergraduate Research Assistant

June 2018 - August 2018

SMALL RNA SEQUENCING (RNA-SEQ) ANALYSIS OF *Drosophila melanogaster loquacious* (LOQS) MUTANTS

- Worked under the supervision of Prof Greg-Tucker Kellogg to write a Snakemake workflow for the analysis of Small RNA-Seq data from *Drosophila melanogaster loqs* mutants.
- Discovered that the presence of a particular Loqs isoform, Loqs-PD is necessary for the biogenesis of hairpin and anti-sense derived Small RNAs.
- The abstract for this work was accepted for the 4<sup>th</sup> RNA Biology Symposium in 2018.

### Honours Project in Computational Biology

January 2019 - December 2019

SCBIMODAL: AN R PACKAGE FOR THE ANALYSIS OF SINGLE CELL RNA-SEQ DATA BASED ON BIMODALITY IN GENE EXPRESSION

- Developed an R package *scbimodal* for the analysis of Single-cell RNA Sequencing data under the supervision of Prof Greg-Tucker Kellogg
- Using Gaussian Mixture Modelling, *scbimodal* identifies genes which exhibit a bimodal distribution and uses their expression values to characterize and cluster cells. The package is currently maintained in a private repository.

## Work Experience

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### Lucence Diagnostics

January 2020 - March 2021

BIOINFORMATICIAN

- Improved and optimised Lucence's flagship *LiquidHALLMARK* diagnostic test for the detection of multiple complex variants.
- Co-invented a patent pending method for the early detection of various cancers using methylated ctDNA (**IPOS App No.10202105843Q**).

### Duke-NUS Medical School (Center for Quantitative Medicine)

July 2021 - July 2023

SENIOR RESEARCH ASSISTANT

- Analyzed large Emergency Department Electronic Health Record datasets using statistical and machine learning methods.
- Validated and fine-tuned several interpretable machine-learning models for the prediction Out-of-Hospital Cardiac Arrest outcomes in Singaporean and German cohorts.
- Provided statistical support for various epidemiological projects relating to emergency medicine.

## Skills

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**Programming Languages** R, Python, Bash, SQL

**Frameworks** Snakemake, Git, LaTeX, Markdown, Docker, AWS

## Selected Publications

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- Xie, F., Zhou, J., Lee, J. W., Tan, M., Li, S., Rajnthern, L. S., ... Liu, N. (2022). Benchmarking emergency department prediction models with machine learning and public electronic health records. *Scientific Data*, 9(1), 1-12.
- Liu, N., Liu, M., Chen, X., Ning, Y., Lee, J. W., Siddiqui, F. J., ... PAROS Clinical Research Network Investigators. (2022). Development and validation of an interpretable prehospital return of spontaneous circulation (P-ROSC) score for patients with out-of-hospital cardiac arrest using machine learning: A retrospective study. *eClinicalMedicine*, 48, 101422.