

# CURRICULUM VITAE

RUNPENG LUO

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EDUCATION	<p><b>Princeton University</b>, Princeton, NJ, USA Doctor of Philosophy (Ph.D.) in Computer Science Sep 2024 - ongoing</p> <p><b>Australian National University</b>, Canberra, ACT, Australia Bachelor of Advanced Computing (Research and Development) (Honours) Feb 2020 - Dec 2023 First Class Honours - GPA: 6.813/7 Advisors: Dr. Yu Lin and Dr. Benjamin Schwessinger</p> <p><b>Strathfield South High School</b>, Sydney, NSW, Australia High School Certificate Oct 2017 - Oct 2019 ATAR: 94.75</p>	
EMPLOYMENT	<p><b>Diversity Arrays Technology</b>, Canberra, ACT, Australia R&amp;D Solution Developer Feb 2024 - Aug 2024</p> <p><b>Australian National University</b>, Canberra, ACT, Australia Technical Assistant at the School of Biology Oct 2022 - Aug 2024 Summer Research Intern at the School of Computing 2021&amp;2022 Summer Teaching Assistant COMP3320 High Performance Scientific Computation 2023 Semester 2 COMP7240 Introduction to Database Concepts 2023 Semester 1 COMP4300/8300 Parallel System 2023 Semester 1 BIOL8002 Advanced Topics in Quantitative Biology and Bioinformatics 2022 Semester 2 COMP2400/6240 Relational Database 2021&amp;2022 Semester 2</p>	
RESEARCH INTEREST	<p><i>Computational Biology</i> <i>Combinatorial Optimization</i> <i>High Performance Computation</i></p>	
RESEARCH EXPERIENCE	<p>Graph Model and Algorithms for Haplotype-resolved Assembly on Dikaryotic Genome Feb 2023 - Dec 2023 Advisors: Dr. Benjamin Schwessinger, Dr. Lianrong Pu, and Dr. Qing Wang Objectives: Design and implement a bi-partition algorithm to assemble and phase dikaryotic genome using third-generation sequencing (TGS) data and Hi-C data.</p> <p>Plasmid Library Diversity Quantification using Nanopore Sequencing Data Jun 2023 - May 2024 Advisors: Dr. Joseph Brock and Dr. Benjamin Schwessinger Objectives: Design and implement a novel classification algorithm to quantify the plasmid combinations from the library mixture.</p> <p>De Novo Reconstruction of Viral Strains via Iterative Path Extraction from Assembly Graphs Dec 2021 - Oct 2022 Advisors: Dr. Yu Lin Objectives: Design and implement an assembly algorithm to reconstruct strains from viral quasispecies under De novo approach.</p>	
AWARDS	<p>ANU Summer Research Scholarship, 2021&amp;2022 Summer, Canberra, Australia NSW Government School International Student Awards - Academic Achievement 2020, Sydney, Australia</p>	
TALKS	<p>27th Annual International Conference on Research in Computational Molecular Biology, (RECOMB 2023), <a href="#">Link</a>, Istanbul, Turkey, Apr 2023.</p>	

RELEVANT SKILLS	<b>Programing:</b> Python, C/C++, Java, Haskell, Rust, ARMv7 Assembly, PostgreSQL <b>Utilities:</b> Bash, L <sup>A</sup> T <sub>E</sub> X, Anaconda, Git, VsCode, Jupyter Notebook
LANGUAGES	<b>Mandarin Chinese</b> (native) <b>English</b> (fluent)
REFERENCES	Dr. Benjamin Schwessinger <i>E-mail:</i> <a href="mailto:benjamin.schwessinger@anu.edu.au">benjamin.schwessinger@anu.edu.au</a> Associate Professor at the Research School of Biology, Australian National University  Dr. Yu Lin <i>E-mail:</i> <a href="mailto:yu.lin@anu.edu.au">yu.lin@anu.edu.au</a> Senior Lecturer at the School of Computing, Australian National University
PUBLICATIONS	Cleaver, A., <b>Luo, R.</b> , Smith, O. B., Murphy, L., Schwessinger, B., and Brock, J., High-throughput optimisation of protein secretion in yeast via an engineered biosensor. <i>Trends in Biotechnology</i> , (2024). <a href="https://doi.org/10.1016/j.tibtech.2024.11.010">10.1016/j.tibtech.2024.11.010</a>  <b>Luo, R.</b> and Lin, Y., VStrains: De Novo Reconstruction of Viral Strains via Iterative Path Extraction From Assembly Graphs. <i>Proceedings of the 27th International Conference in Computational Molecular Biology (RECOMB 2023)</i> , 3-20 (2023). <a href="https://doi.org/10.1007/978-3-031-29119-7_1">10.1007/978-3-031-29119-7_1</a>