Xiang Ge (Xiangge) Luo

□ +41 77 985 76 18 | ■ luox@ethz.ch | ■ xgluo.github.io

Education _____

ETH Zurich Basel, Switzerland

PHD CANDIDATE IN COMPUTATIONAL BIOLOGY

2020.11 - expected 2025.12

- Thesis: Modelling tumor mutation trees for evolution-guided precision oncology
- Supervisor: Prof. Dr. Niko Beerenwinkel | Second advisor: Dr. Jack Kuipers
- Doctoral committee: Prof. Dr. Jasmine Foo, Prof. Dr. Benjamin Raphael

ETH Zurich Zurich. Switzerland

MASTER OF SCIENCE IN STATISTICS

2018.09 - 2020.09

- GPA: 5.88/6 Graduated with distinction
- Thesis: Learning Bayesian networks from ordinal data
- Supervisor: Dr. Jack Kuipers

University of Waterloo Waterloo, Canada 2013.09 - 2018.04 **BACHELOR OF MATHEMATICS**

• GPA: 92% - Graduated with distinction, Dean's Honours List

Research Experience _____

Computational Biology Group, ETH Zurich

Basel, Switzerland

SCIENTIFIC ASSISTANT

2020.11 - present

- · Supervisor: Prof. Dr. Niko Beerenwinkel
- Developed mathematical and statistical models to infer cancer evolution from cross-sectional single-cell phylogenies, aiming to predict drug resistance and facilitate evolution-guided precision oncology.

Institute of Molecular Systems Biology, ETH Zurich

Zurich, Switzerland

SEMESTER STUDENT

2019.03 - 2019.07

- Supervisor: Prof. Dr. Mattia Zampieri
- Analyzed E. coli drug response metabolomics data to investigate regulatory and functional relationships between metabolites.

Seminar for Statistics, ETH Zurich

Zurich, Switzerland

STATISTICS LAB STUDENT

2019.03 - 2019.06

- Supervisor: Dr. Markus Kalisch
- · Analyzed mouse experimental data using mixed models to assess bone loss and identify early osteoporosis markers.

Department of Statistics and Actuarial Science, University of Waterloo

Waterloo, Canada

Undergraduate Researcher

2017.09 - 2017.12

1

- Supervisor: Prof. Dr. Alexander Schied
- Derived the Nash equilibrium for risk-averse investors in a market impact game with transient price impact and quadratic transaction costs over finite and infinite horizons.

Awards & Scholarships ______

2024	SIB Remarkable Outputs 2023 Award, Swiss Institute of Bioinformatics	Switzerland
2017	Undergraduate Student Research Award, NSERC (\$4,500)	Canada
2017	President's Research Award, University of Waterloo (\$1,500)	Canada
2015	President's International Experience Award, University of Waterloo (\$1,500)	Canada
2013	Isabel Farrar Undergraduate Entrance Scholarship, University of Waterloo (\$4,000)	Canada
2013	President's Scholarship of Distinction, University of Waterloo (\$2,000)	Canada

Publications _____

Xiang Ge Luo, Jack Kuipers, Kevin Rupp, Koichi Takahashi and Niko Beerenwinkel. Bayesian inference of fitness landscapes via tree-structured branching processes. Bioinformatics, in press, 2025. (ISMB/ECCB 2025, acceptance rate 17.5%)

Xiang Ge Luo, Jack Kuipers and Niko Beerenwinkel. Joint inference of exclusivity patterns and recurrent trajectories from tumor mutation trees. Nature Communications, 14:3676, 2023. (RECOMB 2022, acceptance rate 21.3%)

Xiang Ge Luo, Giusi Moffa and Jack Kuipers. Learning Bayesian networks from ordinal data. *Journal of Machine Learning Research*, 22:1-44, 2021.

Xiangge Luo and Alexander Schied. Nash equilibrium for risk-averse investors in a market impact game with transient price impact. *Market Microstructure and Liquidity*, 5:2050001, 2020.

Conference	ces, Seminars & Workshops	
2025.07	Contributed talk, ISMB/ECCB 2025 Conference • Bayesian inference of fitness landscapes via tree-structured branching processes	Liverpool, UK
2025.04	 Poster, 6th Zurich Precision Oncology Symposium FiTree: fitness inference from single-cell phylogenies applied to acute myeloid leukemia dat 	Zurich, Switzerland a
2025.03	 Invited talk, D-BSSE Departmental Seminar Series, ETH Zurich Fitness inference from single-cell phylogenies using tree-structured branching processes 	Basel, Switzerland
2025.02	Contributed talk, PhyloBasel Seminar Series, ETH Zurich Bayesian inference of fitness landscapes via tree-structured branching processes	Basel, Switzerland
2024.06	 Award talk, SIB Days – The Swiss Bioinformatics Summit Mining tumor mutation trees with TreeMHN for evolution-guided precision oncology 	Virtual
2023.06	Contributed talk, SKINTEGRITY.CH Annual Retreat • Joint inference of exclusivity patterns and recurrent trajectories from tumor mutation trees	Spiez, Switzerland
2022.06	Contributed talk, Mutual Hazard Networks Workshop • Joint inference of exclusivity patterns and recurrent trajectories from tumor mutation trees	Virtual
2022.05	Contributed talk, RECOMB 2022 Conference • Joint inference of exclusivity patterns and recurrent trajectories from tumor mutation trees	San Diego, USA
2022.04	Contributed talk, OLISSIPO Exchange Week • Joint inference of exclusivity patterns and recurrent trajectories from tumor mutation trees	Lisbon, Portugal
2022.03	Poster, Ascona Workshop 2022 • Joint inference of exclusivity patterns and recurrent trajectories from tumor mutation trees	Ascona, Switzerland
Teaching.		
Spring 2023 Spring 2022 Spring 2022 Spring 2021	Teaching Assistant, Statistical Models in Computational Biology Head Teaching Assistant, Statistical Models in Computational Biology Teaching Assistant, LSZGS Advanced Block Course Computational Biology Teaching Assistant, Statistical Models in Computational Biology	ETH Zurich ETH Zurich ETH Zurich ETH Zurich
Supervision	on	
2023 2023 2022	Laura Quintas, Master Student, INESC-ID (Co-supervised with Monica Baciu-Dragan) Laurenz Keller, Master Student, ETH Zurich (Co-supervised with Pawel Piotr Czyz) Jiayi Wang, Master Student, ETH Zurich (Co-supervised with Pawel Piotr Czyz)	
Academic	Services	
Journal Revi	ewer for Genome Research, Bioinformatics	
Conference R	Reviewer for RECOMB, ISMB	
Language	s and Skills	
Coding	Python, R, C++, MATLAB, Git, Snakemake, ŁTĘX	

Languages English (working proficiency), Mandarin (native), Cantonese (native)