

Lorenzo Bini

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Education

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| University of Geneva , Ph.D. Candidate - Department of Computer Science & CUI | Nov 2022 – Ongoing |
| • Major: Artificial Intelligence and Machine Learning. | |
| • Research interests: Graph neural networks, adversarial learning, representation learning, active and self-supervised learning. Generative AI for medicine and healthcare, including 3D genomics and scRNA data generation. Implementation of robust adversarial models within weak/self-supervised training strategies to reduce label acquisition costs. | |
| Polytechnic of Turin , Master of Science in Physics of Complex Systems | Sep 2020 – Sep 2022 |
| • GPA: 4.0/4.0 | |
| Alma Mater Studiorum - University of Bologna , Bachelor degree in Physics & Astronomy Systems | Sep 2017 – Sep 2020 |
| • GPA: 4.0/4.0 | |

Experience

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| PhD Researcher , Jian Ma's Lab - Comp. Science Dept., Carnegie Mellon University | Sep 2025 - Ongoing |
| • Developed and implemented generative models for epigenomics and gene regulation. | |
| • Designed and integrated LLM-based multi-modal encoders to fuse epigenomic signals and imaging data. | |
| • Implemented scalable training & inference (subchain batching, sparse graphs, RMSD alignment) and evaluation tooling (contact-map, SCC, insulation score, radius-of-gyration). | |
| • Optimized memory and runtime (sparse attention) to enable large-region generation. | |
| PhD Researcher , Hôpitaux Universitaires de Genève (HUG) – Geneva, Switzerland | Nov 2022 - Ongoing |
| • Detection of minimal residual disease (MRD) of acute lymphoblastic and myeloid leukemia (AML/ALL) from flow cytometry data. | |
| • Development of fast training/inference deep learning models (e.g, Graph Transformers) for single-cell classification in weak/self-supervised contexts. | |
| • Development of generative models (e.g., DDPMs, Flow Matching) for 3D genomics, flow cytometry, scRNA-seq, spatial transcriptomic, and multiomics data. | |
| • LLMs to streamline routine hospital processes. | |
| Teaching Assistant , University of Geneva – Department of Computer Science & CUI | Nov 2022 – Ongoing |
| • Introduction to Computational Finance - 14X030 ; | |
| • TALN: Traitement de la langue approches linguistiques et approches empiriques (NLP) - 34C2161 ; | |
| • Information Retrieval - 14X060 ; | |
| • Data Science - Analyse et Traitement de l'Information - 14X026 ; | |
| • Selected Chapters - Game Theory - 14X060 ; | |
| Research Assistant - Intern , Quantum Technology Group - University of Norway | Feb 2022 - July 2022 |
| • Worked on quantum theory for entanglement and non-locality in optomechanics continuous variable systems. Developed quantum frameworks to analyze two-cavity optomechanics systems. | |

Publications

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| LapDDPM: A Conditional Graph Diffusion Model for scRNA-seq Generation with Spectral Adversarial Perturbations | Jun 2025 |
| <i>Lorenzo Bini, Stéphane Marchand-Maillet</i> | |

ICML'2025 + GenBio Workshop: The Second Workshop on Generative AI and Biology, Vancouver.

Self-Supervised Graph Learning via Spectral Bootstrapping and Laplacian-Based Augmentations

May 2025

Lorenzo Bini, Stéphane Marchand-Maillet

Preprint. To appear in 2025, under double-blind review as a conference paper.

Massive Activations in Graph Neural Networks: Decoding Attention for Domain-Dependent Interpretability

Oct 2024

Lorenzo Bini, Marco Sorbi, Stéphane Marchand-Maillet

ECAI'2025, Bologna (oral presentation) + ICLR'2025 Workshop XAI4Science: From Understanding Model Behavior to Discovering New Scientific Knowledge, Singapore.

Injecting Hierarchical Biological Priors into Graph Neural Networks for Flow Cytometry Prediction

Jul 2024

Lorenzo Bini, Stéphane Marchand-Maillet

ICML'2024 + Workshop on Accessible and Efficient Foundation Models for Biological Discovery, Wien, Austria.

FlowCyt: A Comparative Study of Deep Learning Approaches for Multi-Class Classification in Flow Cytometry Benchmarking

Jun 2024

Lorenzo Bini, Margarita Liarou, Thomas Matthes, Stéphane Marchand-Maillet

Conference on Health, Inference, and Learning (CHIL'24), New-York, NY.

Why Attention Graphs Are All We Need: Pioneering Hierarchical Classification of Hematologic Cell Populations with LeukoGraph

Feb 2024

Lorenzo Bini, Thomas Matthes, Stéphane Marchand-Maillet

Preprint arXiv:2402.18610, under double-blind review as a conference paper.

HemaGraph: Breaking Barriers in Hematologic Single Cell Classification with Graph Attention

Dec 2023

Lorenzo Bini, Thomas Matthes, Stéphane Marchand-Maillet

Preprint arXiv:2402.18611, under double-blind review as a conference paper.

Awards & Oral Presentations

Oral Presentation at the Second Workshop on Explainable Artificial Intelligence for the Medical Domain - ECAI'25, Bologna

Oct. 2025

- Invited oral presentation of the accepted paper "Massive Activations in Graph Neural Networks: Decoding Attention for Domain-Dependent Interpretability" at European Conference on Artificial Intelligence (ECAI'25), Bologna.

Winner of Swiss National Science Foundation PhD Mobility Grant

Sep. 2025

- Winner of the Swiss National Science Foundation (SNSF) PhD Mobility Grant for a 6-month internship at Ray and Stephanie Lane Computational Biology Department - Carnegie Mellon University (CMU), Pittsburgh USA.

PhD Symposium - CHIL'24 at Cornell Tech University, NY

Jun. 2024

- Winner of the PhD Symposium money-prize to attend and present PhD work "Adversarial Robust GNNs: Enhancing Learning with Knowledge Injection in Single-Cell Data" at CHIL'24, conference held by Cornell Tech University, New York.

CHAIR Structured Learning Workshop - Chalmers University of Technology

Oct. 2023

- Oral presentation of the "Knowledge Distillation in Acute Myeloid Leukemia Classification: Tabular Data Meets Graph Neural Networks" poster at the AI Structured Learning 2023 Workshop in Göteborg, Sweden.

Winner of Thesis on Proposal 2021/2022

Feb. 2022

- Winner of the "Thesis on Proposal 2021/2022" call for bids for Master's Thesis on "Entanglement and nonlocality in optomechanics continuous variable systems" under the supervision of Prof. Francesco Pietro Massel & Prof. Vittorio Penna.

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- Received grants for research period at USN-Kongsberg.

Projects

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| Flow Cytometry Deep Learning Benchmark | FlowCyt-Benchmark |
| • Developed the first publicly available deep learning benchmark for single cell classification and clustering on flow cytometry data. Tested on a cohort of 30 patients selected by expert hematologists, from bone marrow and peripheral blood samples. Benchmarked SOTA classification/generative models including GNNs, GraphTransformers, Diffusion Models (DDPMs) and VAEs. | |
| • Tools Used: Python, CSS, HTML. | |
| Hackathons & Competitions | 2019 - Ongoing |
| • Regular participation at Kaggle/LeetCode competitions and Hackathons; runner-up BR41N.IO 2021 Hackathon@PoliTO, runner-up UNIBO-IBM-Unipol Hackathon 2019, 3rd classified QuHack4IA 2023. | |
| • Tools Used: Python - Competitive Programming. | |
| Math/Physics Olympiad | Sep 2014 - Sep 2019 |
| • 2x winner of the Italian Regional Math Olympiad. | |
| • 1x winner of the Italian Regional Physics Olympiad. | |
| Visiting Student , City Montessori School, Lucknow - Uttar Pradesh, India USN | Aug 2016 - Sep 2016 |
| • Visited the City Montessori School together with the Italian cultural association "CinemáCinemá" to provide help and needs to elementary/mid school students. | |
| Oxfam Volunteering | 2016 - Ongoing |
| • I do regularly serve as volunteer for charity organization, such as Oxfam Italy . | |

Technologies

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- **Programming Languages:** Python, Julia, R, Matlab, Mathematica, C/C++, CUDA.
 - **Frameworks:** Pytorch, Pytorch-Lightning, Tensorflow, Keras, Jupyter.
 - **AI/ML:** SciPy, scikit-learn, Github, GitLab, Numpy, Pandas, Matplotlib, Seaborn, wandb, PyG.
 - **Tools:** Git, Linux, ssh, SLURM, pip, Anaconda, Docker, VSCode, Bash (Shell).

Software Licenses: Kaluza Analysis Software, Zemax OpticStudio | Comprehensive Optical Design Software, TeXstudio, BioVinci Software.

Reviewing Duties

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| • International Conference on Learning Representations (ICLR). | 2023 - Ongoing |
| • International Conference on Machine Learning (ICML). | 2022 - Ongoing |
| • Learning on Graphs Conference (LOG). | 2023 - Ongoing |
| • AAAI Conference on Artificial Intelligence. | 2025 - Ongoing |
| • IEEE Transactions on Computational Biology and Bioinformatics. | 2023 - Ongoing |