
Lorenzo Bini

Geneva, Switzerland | Lorenzo.Bini@unige.ch | [022 3790124](tel:0223790124) | [Webpage](#) | linkedin.com/in/lorenzo-bini4
github.com/LorenzoBini4

Education

- 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

- 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

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

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.

- Received grants for research period at USN-Kongsberg.

Projects

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

- **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

- 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