Lorenzo Bini

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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

Research Assistant, 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.

Research Intern, Jian Ma's Lab - Comp. Science Dept., Carnegie Mellon University

Sep 2025 - Mar 2026

- 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.

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 Götebor, 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 Compatitive 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 Montessory School, Lucknow - Uttar Pradesh, India USN

Aug 2016 - Sep 2016

• Visited the City Montessory School together with the italian cultural association "CinemíCinemá" 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).	2023 - Ongoing
• Learning on Graphs Conference (LOG).	2023 - Ongoing
AAAI Conference on Artificial Intelligence.	2025 - Ongoing
• IEEE Transactions on Computational Biology and Bioinformatics.	2023 - Ongoing