

Curriculum Vitae Gianmarco Accordi

1 Contacts

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2 Short Bio

I've completed my Master's and Bachelor's degree at Politecnico di Milano, with a thesis titled "A High-throughput pose selection method for extreme scale virtual screening in drug discovery". Currently, I am a PhD student at Politecnico di Milano, and my research thesis is about the acceleration and performance portability of virtual screening applications on emerging HPC architectures. Specifically, I am part of the development team of LiGen, a high-throughput, extreme-scale virtual screening pipeline. LiGen has been used in the largest virtual screening campaign against SARS-CoV-2, the context of the European project EXSCALATE4CoV. In particular, I focus on the porting and optimization of computation kernels on GPUs. My research interests also include the integration of AI modules into virtual screening applications and the analysis of their accuracy and performance.

3 Working Experience

Research Fellow

at Politecnico di Milano 2022 - Present

I'm responsible for designing, implementing, and evaluating techniques to optimize virtual screening applications on modern EuroHPC supercomputers.

Teaching Assistant

at Politecnico di Milano 2022 - Present

I've been a lecturer in various computer science courses on C, Python, and GPU programming.

Full Stack Engineer Specialist

at Mia-Platform 2022

I've worked as a DevOps, focusing on maintaining and developing the ticket-selling system for a railway company. My main contribution has been designing and implementing a comprehensive digital selling system for passes.

Internship

at MobiMESH 2017

The intern's objective was to analyze and design tools and software for a Wireless captive portal for many clients.

4 Education

PhD's student in Computer Science and Engineering
Politecnico di Milano 2022 - Present

Master's degree in Computer Science and Engineering
Politecnico di Milano 2019 - 2022

Bachelor's degree in Computer Science and Engineering
Politecnico di Milano 2016 - 2019

5 Skills

OS Linux, Windows, MacOS

Languages C/C++, Python, Fortran, Bash

Parallel Programming CUDA, OpenMP, OpenACC,
OpenCL, HIP, MPI, SYCL

Compiler NVCC, Clang, GCC, LLVM

AI PyTorch, TensorFlow, Numpy, Pandas, DGL

Cheminformatics LiGen, AutoDock, PyMOL

Project Management Git, Jira

ISA x86, PTX, ARM

Job Scheduler SLURM, PBS

6 Selected Papers

- GPU-optimized Approaches to Molecular Docking-based Virtual Screening in Drug Discovery: A Comparative Analysis*
Submitted to Journal of Parallel and Distributed Computing, 2023
- Out of Kernel Tuning and Optimizations for Portable Large Scale Docking Experiments on GPUs*
Submitted to The Journal of Supercomputing, 2023
- Enabling Performance Portability on the LiGen Drug Discovery Pipeline*
Submitted to Future Generation of Computer Systems, 2023
- Domain-Specific Energy Modeling for Drug Discovery and Magnetohydrodynamics Applications*
Supercomputing, 2023