Lorenzo Di Rocco

☑ lorenzo.dirocco@uniroma1.it

19th of July, 1994

Research Interests

- **Bioinformatics**.
- **High-Performance Computing.**
- Statistical Learning in Computational Genomics and Pangenomics.

Education

2023 – now	Visiting Ph.D Student, Genome Data Science Group , Bielefeld University, Germany. Supervisor: Prof. Dr. Alexander Schönhuth
2021 – now	Ph.D. Methodological Statistics, Sapienza University of Rome, Italy.
2017 – 2020	M.Sc. Statistical and Decision Sciences , Sapienza University of Rome, Italy. Thesis title: <i>Application of Apache Spark for alignment-free analysis of genomic sequences</i> .
2014 - 2017	B.Sc. Economic Sciences, Roma Tre University, Italy.

Academic Positions

2021-2022	Temporary Research Fellow , Sapienza University of Rome, Italy. <i>Start Date</i> : 03/2021 <i>End Date</i> : 02/2022 <i>Supervisor</i> : Prof. Umberto Ferraro Petrillo <i>Description</i> : Winner of a research fellowship granted by the Department of Statistical Sciences at Sapienza University of Rome.
2020-2021	Research Project Associate, Sapienza University of Rome, Italy.Start Date: 08/2020End Date: 01/2021Supervisor: Prof. Umberto Ferraro PetrilloDescription: I collaborated with the Department of Statistical Sciences at Sapienza - University of Rome, at the design and the development of distributed algorithms for the analysisof protein-to-protein interaction networks, under the framework of the EIT Health European project on "Big Data Software Services for Decision Support in Precision Medicine".

Summer Schools

5-8 July 2022	Introduction to Pangenomics, Lake Como School of Advances Studies, Italy. Or-
	ganizers: Prof. Paola Bonizzoni (Univ. Milano-Bicocca),Prof. Gianluca Della Vedova
	(Univ. Milano-Bicocca), Prof. Alex Schoenhuth (Univ. Bielefeld, Germany)
3-15 July 2023	Summer School on Parallel Computing, CINECA - Bologna Offices Italy.
	Organizers: Dr. Emerson Andrew (CINECA).



Projects

Active Projects

Principale Investigator of a Class C ISCRA Project

Title: Efficient Pangenome-based Inference for Genomic Variants using New Era Statistical Methods.

Value: 20,000 CORE/h to spend on Galileo100 system at CINECA

- Member of Research Project funded by Sapienza University of Rome *Title*: Advancing Precision Public Health Medicine: High-Performance Computing for Efficient Solutions to Bioinformatics and Biostatistical Challenges
- Member of Research Project funded by Sapienza University of Rome Title: Unlocking the Potential of Artificial Intelligence for Generating Synthetic Patients in Clinical Trials

Past Projects

2021-2023

Collaborator on the AIGREET Project

*Supervisor:*Prof. Umberto Ferraro Petrillo *Description:* In partnership with Sapienza University of Rome, Andersen Italia and Internationalia, the AIGreet project exploits web scraping techniques and artificial intelligence to automatically recognize and track the progress of green investments.

Events

2024 CIBB 2024 - 19TH Conference on Computational Intelligence Methods For Bioinformatics and Biostatistics
Pela Decempite Merchan

Role: Program Committe Member

2023 CIBB 2023 - 18TH Conference on Computational Intelligence Methods For Bioinformatics and Biostatistics *Role:* Program Committe Member

Special Session in Distributed Computing in Bioinformatics and Computational Biology *Role:* Organizer

Teaching Experience

 2022 - 2023 Introduction to Deep Learning Role: Teaching Assistant Institution: Specialization School in Health Statistics and Biometry.
 Introduction to Statistical Learning

> *Role:* Teaching Assistant *Institution:* Department of Statistical Sciences, Sapienza University of Rome.

Teaching Experience (continued)

2021 - 2022

R-Programming

Role: Teacher

Institution: Advanced Training Course in Bioinformatics, Sapienza University of Rome.

Big Data Management

Role: Teaching Assistant *Institution:* Department of Statistical Sciences, Sapienza University of Rome.

Skills

Coding Python, C++, Java, Scala, R Databases MysQL, Neo4J, MongoDB.

Research Publications

- L. Di Rocco and U. Ferraro Petrillo, "A distributed alignment-free pipeline for human snps genotyping," in Proceedings of the 14th ACM International Conference on Bioinformatics, Computational Biology, and Health Informatics, 2023, pp. 1–8.
- L. Di Rocco, U. F. Petrillo, and G. Grani, "Enhancing scalability of distributed snps calling pipelines using cluster-driven partitioning strategy," 2023.

G. Grani, L. Di Rocco, and U. F. Petrillo, "Using knowledge graphs to model green investment opportunities," in *European Conference on Advances in Databases and Information Systems*, Springer, 2023, pp. 440–451.

L. Amorosi, L. D. Rocco, and U. F. Petrillo, "Scheduling k-mers counting in a distributed environment," in Optimization in Artificial Intelligence and Data Sciences: ODS, First Hybrid Conference, Rome, Italy, September 14-17, 2021, Springer, 2022, pp. 73–83.

L. Di Rocco, U. Ferraro Petrillo, and S. E. Rombo, "Diamin: A software library for the distributed analysis of large-scale molecular interaction networks," *BMC bioinformatics*, vol. 23, no. 1, p. 474, 2022.

L. Di Rocco, U. F. Petrillo, and F. Palini, "Large scale graph based network forensics analysis," in *Pattern Recognition. ICPR International Workshops and Challenges: Virtual Event, January 10–15, 2021, Proceedings, Part V*, Springer, 2021, pp. 457–469.