

# Dany Mukesha

*PhD Candidate in Bioinformatics*

## Education

- 2023–2026 **Doctor of Philosophy, Bioinformatics**, *Université Côte d'Azur*, Nice, France  
2020–2023 **Master's degree, Bioinformatics**, *University of Rome Tor Vergata*, Rome, Italy  
2016–2019 **Bachelor's degree, Bioinformatics**, *Sapienza Università di Roma*, Rome, Italy

## Experience

- March 2022 – **Bioinformatician**, *Wayne State University*, Detroit, Michigan, United States  
Sept. 2023 Performed computational analysis for gene expression data to evaluate the effects of environmental contaminants, such as plasticizers, on cardiovascular risk and disease.
- Dec. 2021 – **Software Engineer**, *Accenture SpA*, Rome, Latium, Italy  
Jan. 2023 Supported the implementation of software solutions for companies to conduct business operations and processes at enterprise-level.
- Sept. 2020 – **Bioinformatician**, *The University of Manchester*, Manchester, England, UK  
August 2021 Integrated epigenetic data and GWAS results to identify causal variants, genes, and cell types in Rheumatoid Arthritis (RA). Updated credible SNPs list through analysis of unrelated samples, assuming linkage disequilibrium between marker alleles and disease susceptibility alleles, improving RA genetic risk mapping.
- Jan. 2021 – **TIBCO Developer**, *Catenate Group*, Rome, Latium, Italy  
Dec. 2021 Developed and maintained TIBCO applications, including setting up Mashery API with security protocols. Created BusinessWorks processes for UAT and monitored TIBCO EAI applications for billing and inventory synchronization.
- July 2020 – **Bioinformatician**, *Policlinico Agostino Gemelli IRCCS*, Rome, Latium, Italy  
Dec. 2020 Performed transcriptomic analysis of ependymoma tumor samples, identifying Hub genes and key pathways through gene co-expression network analysis. Findings published in the International Journal of Molecular Sciences.
- Jan. 2019 – **Bioinformatician**, *Sapienza Università di Roma*, Rome, Latium, Italy  
Oct. 2019 Conducted genomic analysis of Burkholderia cepacia complex (BCC) isolates to identify genetic mutations and classify phylogenetic subgroups, revealing distinct virulence factors and antibiotic resistance profiles associated with each subgroup.

## Publications

- [Mukesha, 2024] Mukesha, D. Scalable and interpretable genetic algorithm framework for high-throughput genomic data. 2024. doi:10.5281/zenodo.15801072.
- [Mukesha et al., 2025] Mukesha, D., Sarter, M., Dubray, M., et al. Targeted serum metabolomic profiling and machine learning approach in alzheimer's disease using the alzheimer's disease diagnostics clinical study (addia) cohort. *medRxiv*, 2025. doi:10.1101/2025.04.01.25325021.

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## Software Contributions

- 2025 **BioGA**: Bioinformatics Genetic Algorithm R package (v0.99.16).  
<https://bioconductor.org/packages/BioGA/>

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## Other Publications

- 2021 Servidei T, Meco D, Martini M, et al. *The BET Inhibitor OTX015 Exhibits In Vitro and In Vivo Antitumor Activity in Pediatric Ependymoma Stem Cell Models*. **Int J Mol Sci**. 2021;22(4):1877. <https://doi.org/10.3390/ijms22041877>  
Contribution: Performed correlation network analysis of large, high-dimensional datasets.

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## Conferences and Awards

- April 1–5, 2025 Poster-Onsite at **AD/PD™ 2025** International Conference, Vienna, Austria.  
Topic: Metabolomic Signatures and Machine Learning Models for Distinguishing Alzheimer's Disease and Dementia with Lewy Bodies.
- March 5–9, 2024 Poster-Onsite at **AD/PD™ 2024** International Conference, Lisbon, Portugal.  
Topic: Updated panel of genetic variants associated with multiple forms of neuro-degenerative diseases.
- 2023–2026 **PhD Scholarship funded by European Union (EU) program** (France):  
Marie Skłodowska-Curie Actions (MSCA) Doctoral Network.
- 2016–2022 **LazioDisco Scholarship of Merit** (Italy):  
Awarded for academic excellence during Bachelor's (2016–2019) and Master's (2020–2022) studies.

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## Skills

<b>Bioinformatics</b>	Genomics, transcriptomics, GWAS, gene co-expression network analysis
<b>Statistics/ML</b>	Regression (Linear, Logistic, Mixed), Clustering (K-means, HCA), Neural Networks, Gaussian processes, Feature and Model Selection, PCA/ICA, LDA, KNN, RF, Bagging, Bayesian methods, CV, Meta-analysis, Text mining.
<b>Software-Dev.</b>	Java microservices, Spring Boot, REST APIs, Angular, React
<b>DevOps &amp; Tools</b>	Kubernetes, Linux, Bash scripting, network configuration, Git, Docker, Terraform, Jenkins, GitLab/GitHub, LDAP.
<b>Operations</b>	Unix, Cloud, Containers
<b>Databases</b>	Oracle, PostgreSQL, MySQL
<b>Languages</b>	English, Italian, French
<b>Programming</b>	C++, Python/R, Java, Bash

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