Olusegun Ekundayo Adebayo

olusegun.adebayo@college-de-france.fr

Work Experience

Bioinformatician/Post-doc Researcher

Collège de France, CIRB - Denis Duboule

09/2025 - current

- Conducting research on **chromatin organization and developmental gene regulation** using large-scale 2D/3D omics multi-omics datasets.
- Administering and maintaining a **local Galaxy server** to provide reproducible, user-friendly bioinformatics workflows for lab members.
- Applying **DevOps practices** (Ansible for automation, Singularity/Docker for containerized workflows, Git/GitHub for version control) to ensure maintainable and scalable computational environments.
- Developing and integrating **custom workflows** into Galaxy for seamless data analysis.
- Performing bioinformatics analyses and visualization in **Python**, **R**, and **Bash**, translating computational results into biological insights and publications.
- Supporting collaborative research by ensuring workflow reproducibility, scalability, and security.

Bioinformatician/Post-doc Researcher

05/2025 - 08/2025

University of L'Aquila, DISIM - Antinisca Di Marco

- Developed pipelines to interpret BRCA1 and BRCA2 variants in cDNA format and generate mutated FASTA sequences based on CDS references.
- Implemented algorithms to inject **combinations of pathogenic and non-pathogenic mutations** while handling positional shifts to maintain sequence integrity.
- Created automated workflows for window-based mutation extraction (500–1000 bp) to support downstream analysis on HPC clusters.
- Applied machine learning classification using DNABERT2 to predict variant effects, integrating results into analysis pipelines.
- Maintained reproducibility and transparency with **Python pipelines** and version-controlled code hosted on **GitHub**.

PhD Researcher 03/2022 - 03/2025

Université Marie & Louis Pasteur, LmB - Raluca Eftimie

- Computer Vision with Python, OpenCV and YOLO.
- Conduct data analysis using Python (Pandas, NumPy) for biological and bio-mechanical experiments.
- Develop machine learning models using **Keras**, **Tensorflow**, and **scikit-learn** and deploy them using FastAPI and Streamlit.
- Utilize **FEniCS** for finite element analysis.
- Present research results at international conferences and collaborate with interdisciplinary teams.
- Develop local and non-local mechanistic mathematical models to describe biological and medical problems.
- Use Git and Github to store and host codes.

Data Scientist

10/2021 - 02/2022

- Livingston Research, Rome, Italy
 - Applied data analysis and machine learning algorithms in R and Python to solve client problems.
 - \bullet Delivered content related to ${\bf statistical}$ analysis and ${\bf machine}$ learning concepts.
 - Used advanced Excel, Matplotlib and seaborn for data analysis and visualization

PhD in Applied Mathematics. Université de Bourgogne Franche-Comté (UBFC).	03/2022 - 03/2025
Master degree in Artificial intelligence and Machine learning Simplilearn Solutions	08/2021 - 03/2024
Joint Master degree in Mathematical Modeling In Engineering Technische Universität Wien, Vienna (Austria) Mathmods.	09/2019 - 09/2021
Joint Master degree in Mathematical Modelling In Engineering Università degli Studi dell'Aquila. (Italy) Mathmods.	09/2019 - 09/2021
Bachelor of Science BSc. Mathematics, Obafemi Awolowo University. Ile-Ife (Nigeria).	06/2014 - 02/2018

Digital Skills

- Data Analysis & Machine Learning: Python scikit-learn, Keras, Tensorflow, PyTorch.
- Workflow & Automation: Snakemake, Nextflow, runHiC, Ansible, Git/GitHub
- Containers: Singularity, Docker
- Bioinformatics: Galaxy server administration, Hi-C, RNA-seq, multi-omics analysis, HPC
- Visualization: Excel, R, Matplotlib, Seaborn, PowerBI, Paraview, ggplot2, Plotly, Dash.
- Model Deployment: Streamlit, Tensorflow Server, FastAPI
- Programming Languages: Bash, Python, MATLAB, R, C/C++
- Operating Systems: Linux/Unix, Windows, MacOS, SLURM cluster server.
- Database SQL.
- Cloud Services GCP, AWS.

Honor and Awards

AfriGen-D IBT_2025 – Introduction to Bioinformatics Training. University of Cape Town, AfriGen-D [2025]: Selected participant in this competitive hybrid/online bioinformatics training course.

Financial support towards the XLIV Dynamics Days Europe (DDE'24 – Bremen), Constructor University, Bremen, Germany [2024]: Funded by the wilhelm und else heraeus-stiftung

International Doctoral student Mobility to the University of Dundee, Scotland [2024]: Funded by Université de Bourgogne Franche-Comté ("projet mobilité internationale 2024")

PhD scholarship in Applied Mathematics from the Université de Bourgogne Franche-Comté (UBFC) [2022]: Funded by ANR-PRCI Project.

Mathmods Consortium Thesis Grant [2022]: The award was given to those who completed their thesis at the appropriate time

MathMods Consortium Scholarship. Mathmods Consortium (the University of L'Aquila, TU Vienna, University of Hamburg, University of Cote d'Azur)[2019-2021]: The award covered the first and second year of my Masters thesis. Azienda per il Diritto agli Studi Universitari L'Aquila (ADSU) Scholarship

University of L'Aquila Grant [2019-2020]: The award covered the first year of my Masters thesis and was awarded to the top 20 on the ranking list.

Publications

Adebayo, O. E., Trucu, D., and Eftimie, R. (2025c). Condition number for finite element discretisation of nonlocal PDE systems with applications to biology. arXiv preprint (2025) arXiv:2508.09781

Adebayo, O.E., Chatelain, B., Eftimie, R., Trucu, D. Deep learning approaches for classifying images of keloids in the context of malignant and benign skin disorders, *Diagnostics*, **15** (6)(2025), 710

Adebayo, O.E., Trucu, D., Eftimie, R., Analytical investigation of a non-local mathematical model for normal and abnormal wound healing. *Discrete and Continuous Dynamical Systems - B.*, **30** (7) (2024) 2401-2428

Adebayo, O.E., Urcun, S., Rolin, G., Bordas, S., Trucu, D., Eftimie, R.: Mathematical investigation of normal and abnormal wound healing dynamics: local and non-local models. *Mathematical Biosciences and Engineering*, 20, (2023), 17446 - 17498

Eftimie, R., Rolin, G., **Adebayo**, **O.E.**, Urcun, S., Chouly, F., Bordas, S.: Modelling keloid dynamics: a brief review and new mathematical perspectives. *Bulletin of Mathematical Biology*, **85** (2023), 117

Certifications

\bullet Introduction to Bioinformatics: AfriGen-D $\parallel University$ of Cape-Town	08/2025
• Artificial Intelligence Engineer: Simplilearn	04/2024
• Deep Learning with Keras and Tensorflow: Simplilearn	09/2023
• Machine Learning: Simplilearn	09/2023
• Data Science with Python: Simplifearn	09/2023
• Introduction to Artificial Intelligence: Simplifearn	09/2021
$ \bullet \ \ \text{Applied Plotting, Charting \& Data Representation in Python: } \ \textit{Coursera} \ \textit{University of Michigan} \\$	10/2020
\bullet Data Analysis with Python: Coursera $\ IBM$	09/2020
• Introduction to HTML5: Coursera University of Michigan	03/2019

Presentations

XLIV Dynamics Days Europe (DDE'24 – Bremen, Constructor University). Investigating normal and abnormal wound healing: from deterministic nonlocal mathematical models to machine learning [July-August 2024].

2nd SSEN Annual Postgraduate Research Symposium (University of Dundee). Modeling and Analysis of Non-local Interactions in Wound Healing: Numerical Simulations, Analytical Investigations, and Machine-Learning Approaches [May 2024].

Journées numériques de Besançon 2024: Data driven computing and modeling in biology. *Mathematical modeling of wound healing: local and non-local models* [January 2024].

XXIIeme Journee de l'ED Carnot Pasteur Universite de Franche-Comte. Mathematical modeling of wound healing: local and non-local [June 2023].