Sarani NATIVEL-SANTERNE

Bioinformatician

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EDUCATION

Université de Lille

Masters in Bioinformatics; major: 15.44/20

Université de Lille

Bachelors in Biology of Organisms and Populations

Villeneuve d'Ascq, Lille, France Sept. 2023 – Sept. 2025 Villeneuve d'Ascq, Lille, France Sept. 2020 – May 2023

SKILLS

Tech Stack: Python, R, LaTex, Bash, Bioinformatics tools (BLAST, QIIME, etc.), Biostatistics (Survival analysis, Differential RNA-seq analysis, Discriminatory analysis, PCA, Clustering), Machine Learning

Transferable Skills: : Research, Autonomy, Curiosity, Adaptability, Patience

Languages: : French (Native), Reunionese Creole (Native), English (Bilingual), Spanish (Professional Working Proficiency), Japanese (Limited Working Proficiency), Hindi (Elementary Proficiency), Latin, notions of Odia

EXPERIENCE

GD Biotech

Bioinformatician (apprenticeship)

Lille, France Sept. 2024 – Present

- Research: Pursued a research project about the Eukaryome. Analyzed raw 18S sequencing data.
- Industry: Analyzed microbiome data using R packages (phyloseq, microbiome, microbiomeutilities). Computed statistics and diversity metrics using QIIME.

Université de Lille

 $Bioinformatician\ (internship)$

Villeneuve d'Ascq, Lille, France Aug 2023 – Mar 2024

• Variants Analysis: Analyzed the presence of insertions in the genome of *Arabidopsis halleri* using Python and R. Created a scientific poster to summarize findings.

CERTIFICATIONS

Online courses: Completed several online courses (certifications available here).

Diplomas: Baccalaureate (2019, with highest honors), Bachelor's degree (2023, with honors)

Projects

Participation to the HackBio online internship: A 5-stages online internship exploring different aspects of bioinformatics. All codes and productions are available here, as well as the certificate of participation. Developed a website to summarize the work done at HackBio for a university class.

Platypus Genome Literature Review and bioinformatics analysis: Currently writing a literature review about the platypus' genome and re-doing past bioinformatics analysis on the latest platypus genome.