# 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

Lille, France

Sept. 2024 - Present

 $Bioinformatician\ (apprenticeship)$ 

- 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 and re-doing past bioinformatics analysis on the latest platypus genome.