



- » **Status:** PhD student in Bioinformatics and Statistics applied to plant biology under climate change, BPMP team, Supagro, Montpellier
- » **Fields:** Genomics, Epigenomics, Gene Regulatory Networks, Statistics, Machine Learning, Computational Biology.

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»» Education

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|-------------|---|--------------------------------------|
| 2019 - 2022 | PhD project: Statistical inference of the gene regulatory network in <i>A. thaliana</i> under climate change | CNRS - BPMP & IMAG
research units |
| | <ul style="list-style-type: none"> » A majority of cultivated plants exposed to elevated CO2 undergo a depletion of their nutritional content. My project is to integrate dynamic transcriptomes and epigenomes and from <i>A. thaliana</i> plants to understand how elevated CO2 is linked to mineral nutrition in plants, using statistical learning and predictive models. » Supervisors : Dr Antoine Martin and Dr Sophie Lebre. | |
| 2018 - 2019 | MASTER degree in Artificial Intelligence | University of Lyon 1 |
| | <ul style="list-style-type: none"> » Courses covering machine learning, deep learning, multi-agents systems, reinforcement learning, bio-inspired intelligence, data visualization, AI and cognition. » Double degree in partnership with the INSA (National Institute of Applied Sciences, Lyon). | |
| 2014 - 2019 | Engineering degree in Bioinformatics and Modelling | INSA, Lyon |
| | <ul style="list-style-type: none"> » 2016 - 2019 : Informatics, statistics, algebra, differential equations for modelling and analysis of biological mechanisms. » 2014 - 2016 : 2-years preparatory classes in european section : general scientific and engineering training. | |
| 2014 | Baccalaureate certificate | Ain, France |
| | <ul style="list-style-type: none"> » Engineering Sciences diploma with high honours. Specialization : Mathematics | |

»» Work experience

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| February-July
2019 | Master internship : Machine learning models to predict gene regulation in different cell types | IGMM (CNRS)-LIRMM,
Montpellier |
| | <ul style="list-style-type: none"> » Classification of enhancer-gene interactions in the context of regulation via chromatin folding. » Comparative study of penalized logistic regression, ensemble methods (Random Forest, Boosting), and deep learning models for this task. Languages : R, Python, bash | |

May-July 2018	Internship : tracking algorithm for 3D biological objects	Nanolive, Lausanne
	» Creation of image analysis tools for quantitative biology, inside the Nanolive R&D team. Segmentation of objects and tracking methods for biological objects in label-free microscopy images. Programming language : Python.	
Juillet 2017	Research internship in bioinformatics	LIRMM, Montpellier
	» Enhancement of a supervised classification method to predict transcription factors binding sites. Involved in a scientific paper in BMC Genomics (Link here). Programming language : R.	
2016-2017	Project : model of cancerous cell mutations	NovaDiscovery, Lyon
	» Realized a project in partnership with NovaDiscovery, a consulting Start up in Bioinformatic. Implementation of an object oriented model for mutation dynamics in carcinogenesis.	
2016-2017	Teaching experiences	INSA, Lyon
	» Tutoring a group of four students of INSA preparatory classes in Mathematics. Particular lessons to high school students in mathematics, physics, chemistry	
July 2015	Working internship abroad	Sercotel Espana
	» Waitress in a Catalan hotel.	

»» Skills

Informatics

- » Programming languages : R (package development, Rmarkdown, R-Shiny), Python, Latex, Java, bash, C++, SQL
- » Logiciels : Git, Microsoft Office, Jade, Knime

Interpersonal skills

- » Spoken languages : fluent english (TOEIC) and Spanish (DELE B2)
- » Ability to listen, work in group, adapt, work autonomously
- » First aid diploma PSC1, driving licence B et A2
- » Interests : sports, drawing