# ANTOINE BICHAT

#### Mathematical Engineer - PhD in Biostatistics

antoine.bichat@protonmail.com

### PROFESSIONAL EXPERIENCE

Today   2021	•	Computational scientist Servier Paris area, France • Disease understanding, statistical modeling • Oncology, hard-to-treat cancers, pediatric cancers	
2020   2018	•	Industrial PhD StudentEnterome♥ Paris, France• PhD with CIFRE convention	
2017	•	Biostatistician intern Nestlé Research Center♥ Lausanne, Switzerland• Data analysis of microbiome 16S sequencing • Implementation of DADA2 pipeline for sample inference	IN ଜ ଡ
2016	•	Biostatistician intern Enterome ♀ Paris, France • Data analysis of microbiome Illumina next-generation sequencing • Writing and automation of bioinformatics quality control report	in Ƴ Q
		EDUCATION	
2020   2018	•	<ul> <li>PhD</li> <li>Université Paris-Saclay</li> <li>Discovering multi-scale metagenomic signatures through hierarchical organization of species</li> <li>Tutored by Dr Christophe Ambroise, Dr Mahendra Mariadassou and Mr Jonathan Plassais</li> </ul>	
2017   2016	•	Mathematics of Modelling Master's degreeSorbonne Université• Applied Mathematics for Biology and Medical Sciences• Data Science	
2017   2014	•	Graduate Engineering School Mines Nancy ● Mathematical engineering ● Bioengineering	TE Hig vis Ski
	۲	Classe Préparatoire aux Grandes ÉcolesLycée du Parc, MP*♥ Lyon, France	an



#### LANGUAGES

- French (mother tongue)
- English (fluent C1, IELTS)
- Italian (proficient B2, CILS)

#### TECHNICAL SKILLS

Highly skilled in R, tidyverse and visualization

Skilled in Python, Matlab, LaTeX and Office

Knowledge in C++, SQL, HTML and CSS

## TEACHING EXPERIENCE

Today   2019	•	<b>Data Science Starter Program</b> École Polytechnique, 21h, continuing education ♀ Paris area, France
2020   2018	•	An Introduction to the Linear ModelAgroParisTech, 40h, engineering students♥ Paris, France
2017	•	Visualization with R Mines Nancy, 3h, engineering students ♀ Nancy, France
	2	SCIENTIFIC PRODUCTIONS
2020	•	Hierarchical correction of p-values via a tree running Ornstein-Uhlenbeck process Preprint, submited to <i>Computational Statistics</i>
2020	•	Incorporating phylogenetic information in microbiome abundance studies has no effect on detection power and FDR control Published in <i>Frontiers in Microbiology</i>
2019	•	Quantifying the impact of tree choice in metagenomics         differential abundance studies with R         Poster for UseR! 2019         • Winner of Biostatistics poster prize
2019	•	Impact of tree choice in metagenomics differential abundance studies Talk at SPMGD & JdS Parcelona, Spain & Nancy, France
		VOLUNTEER WORK
2021   2018		<ul> <li></li></ul>
2017   2014		During Engineering School Mines Nancy♥ Nancy, France• Academic year-group representative • Student member of the Executive Board of Mines Nancy • President of the debating and conference club • Communication Officer for the recruitment forum of Mines Nancy
2014   2011	•	UDPS 69 (first aid association)First aid worker, lifeguardQ Lyon area, France