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Jasmine Gamblin

Education

2020 – present PhD in population biology, SORBONNE UNIVERSITÉ, Paris, France

Modelling and inference for the evolution of *Eschrichia coli* accessory genome. Supervised by Amaury Lambert and François Blanquart from the team Stochastic Models for the Inference of Life Evolution (CIRB, Collège de France).

- 2018 2020 **Computational Biology and Bioinformatics Master**, ETH, Zurich, Switzerland Mathematical models for evolution, genetics, data analysis (GPA 5.5/6).
- 2015 2018 **Engineering Degree in Bioinformatics**, ÉCOLE POLYTECHNIQUE, Palaiseau, France Interdisciplinary cursus in computer science, biology and mathematics (ranked 96/428, GPA 3.83/4).
- 2013 2015 **Mathematics and Physics Preparatory Class**, LYCÉE HENRI IV, Paris, France Two year intensive training in mathematics, physics and computer science to prepare the competitive exam entrance to the French *Grandes Ecoles*.

Research Internships

2020 Master's thesis, 6 months, Collège de France, Paris, France

Modelling the dynamics of phage adaptation to a complex environment using branching processes. Supervised by Amaury Lambert and François Blanquart, team Stochastic Models for the Inference of Life Evolution.

2019 Lab rotation, 6 weeks, ETH, Zurich, Switzerland

Applied a new computational method inferring gene regulation networks from RNA expression data in order to compare human and mouse kidney cells. Supervised by Pr. Joshua Payne, Computational Biology group.

- 2019 Lab rotation, 6 weeks, ETH, Basel, Switzerland
 - Implemented an R package allowing to simulate phylogenetic trees under a Birth-Death model conditioned on sampling times. Supervised by Dr. Timothy Vaughan, Computation Evolution group.
- 2018 Research Intern, 4 months, McGill University, Montréal, Canada

Developed an algorithm allowing to adapt BLAST to probabilistic genomes alignment. Supervised by Pr. Mathieu Blanchette, Computational Genomics lab.

2017 **R&D Intern, 2 months**, STILLA TECHNOLOGIES, Villejuif (FR)

Studied the kinetics of the PCR (Polymerase Chain Reaction) in the context of the microfluidic devices produced by Stilla.

Teaching

2020 – 2023 **Teaching Assistant in Mathematics**, SORBONNE UNIVERSITÉ, Paris, France

Undergrad level:

- Mathematics for science (2 groups, 1 semester)
- Introduction to probability theory (1 semester)
- Python for mathematics (2 semesters)
- Oral interrogations in mathematics (1 semester)

2015 – 2016 **Education Assistant**, APPRENTIS D'AUTEUIL, Nantes, France Supervised the daily life of teenagers in a boarding school. Organized educational activities, taught mathematics, English and French during extra tutoring.

Publications

- Jasmine Gamblin, Sylvain Gandon, François Blanquart, Amaury Lambert, Bottlenecks can constrain and channel evolutionary paths. *Genetics, in press* (2023). doi: 10.1093/genetics/iyad001
- Félix Foutel-Rodier, François Blanquart, Philibert Courau, Peter Czuppon, Jean-Jil Duchamps, Jasmine Gamblin, Élise Kerdoncuff, Rob Kulathinal, Léo Régnier, Laura Vuduc, Amaury Lambert, Emmanuel Schertzer, From individual-based epidemic model to McKendrick-von Foerster PDEs: a guide to modeling and inferring COVID-19 dynamics. *Journal of Mathematical Biology* 85, 43 (2022). doi: 10.1007/s00285-022-01794-4

Presentations

Invited talks

- Nov. 2022 Seminar of the Evolutionary Theory department (Max Planck Institute for Evolutionary Biology), Plön, Germany
- Sep. 2022 Workshop evolution and phages, Montpellier, France

Contributed talks

- Jan. 2023 Interdisciplinary Approaches in Molecular Evolution (AIEM), Grenoble, France
- Jan. 2023 PopGroup56, London, UK
- Aug. 2022 Congress of the European Society for Evolutionary Biology (ESEB), Prague, Czech Republic

Posters

- Jun. 2022 Mathematical and Computational Evolutionary Biology (MCEB), Château d'Oex, Switzerland
- Mar. 2022 Interdisciplinary Approaches in Molecular Evolution (AIEM), Rennes, France

Others

Programming PYTHON, R (advanced), C++, JAVA, OCAML (intermediate)

Languages French (Mothertongue), English (Fluent), German (B1)

Organization Abstract selection for the French Natural History Museum young reasearchers' Congress