Morjani Lab, University of California, Berkeley Stanley Hall, Rm 324D Berkeley, CA 94720-3220 ℘ (+1) 341 766 8054 ⋈ kerdoncuff@berkeley.edu ≅ ekerdoncuff.github.io

Élise Kerdoncuff

Education and Professional Experience

- 2022-current **Postdoc**, Moorjani Lab at University of California, Berkeley, USA. Study of South Asian population evolutionary history and development of local ancestry inference methods.
 - 2017-2021 PhD, Detection of Recent Decline in Population by a Genomic Approach, Stochastic Models for the Inference of Life Evolution group (CIRB, Collège de France) and Atelier de Bio-Informatique group (Muséum National d'Histoire Naturelle - MNHN). Supervised by Guillaume Achaz and Amaury Lambert, Sorbonne University.
 - 2015-2017 Master, Ecology, Biodiversity, Evolution, speciality: Evolutionary Ecology, Sorbonne University.
 - 2012-2015 **Bachelor**, Bi-disciplinary bachelor of biology and mathematics, Sorbonne University at the Roscoff Marine Station, France.

Exchange, 3^{rd} year of bachelor (09/2014-06/2015) at the National University of Singapore, Singapore.

Publications, Preprints

- 1. **Kerdoncuff, E.***, Skov L.*, Patterson N., Zhao W., Lueng Y.K., Schellenberg G.D., Smith J., Ganna A., Dey AB., Kardia S., Lee J., Moorjani P. 50,000 years of Evolutionary History of India: Insights from 2,700 Whole Genome Sequences. *bioRxiv* preprint (*EK and LS contributed equally to this work)
- 2. **Kerdoncuff, E.**, Lambert, A., Achaz, G. How demography shapes linkage disequilibrium with or without recombination. *bioRxiv* preprint
- 3. Freund, F.*, **Kerdoncuff, E.***, Matuszewski, S., Lapierre, M., Hildebrandt, M., Jensen, J. D., Ferretti, L., Lambert, A., Sackton, T. B., Achaz, G. Interpreting the pervasive observation of U-shaped Site Frequency Spectra. (2023) *PLOS Genetics* (*FF and EK contributed equally to this work)
- Foutel-Rodier, F., Blanquart, F., Courau, P., Czuppon, P., Duchamps, J., Gamblin, J., Kerdoncuff, E., Kulathinal, R., Régnier, L., Vuduc, L., Lambert, A., Schertzer, E. From individual-based epidemic model to McKendrick-von Foerster PDEs: a guide to modeling and inferring COVID-19 dynamics. (2022) *Journal of Mathematical Biology*
- 5. **Kerdoncuff, E.**, Lambert, A., Achaz, G. Testing for population decline using maximal linkage disequilibrium blocks. (2020) *Theoretical Population Biology.*
- Ribardière, A., Centanni, J., Dano, A., Coudret, J., Daguin-Thiébaut, C. Houbin, C., Kerdoncuff, E., Jambut, S., Cordaux, R., Broquet, T. Female-biased sex ratios unrelated to Wolbachia infection in European species of the Jaera albifrons complex (marine isopods). (2018) Journal of Experimental Marine Biology and Ecology.

Presentations

Invited

- o Human Evolutionary Genetics group weekly seminar, Mai 2024, Institut Pasteur, Paris, France.
- Workshop on Li-Stephens based methods and applications, December 2022, Harvard, USA.

- Center for Theoretical and Evolutionary Genetics weekly seminar, April 2022, UC Berkeley, USA.
- Workshop: Evolutionary Genomics and Life History Traits, October 2019, Montpellier, France.
- o Séminaire du CESCO, June 2019, MNHN, Paris, France.

Contributed talks

- New Frontiers in Reconstructing Human Evolution History, June 2023, Hannover, Germany.
- o Bay Area Population Genomics, October 2022, Berkeley, USA.
- o Popgroup53, January 2020, Leicester, Royaume-Uni.
- o GDR Approches Interdisciplinaires en Evolution Moléculaire (AIEM), November 2019, Toulouse, France.
- Colloque annuel Society for Molecular Biology and Evolution (SMBE), Symposium: The Coalescent in the Era of Population-Scale Genomics, July 2019, Manchester, Royaume-Uni.
- Mathematical and Computational Evolutionary Biology (MCEB), May 2019, Porquerolles, France.
- o Séminaire de l'ISYEB, May 2019, MNHN, Paris, France.
- o CIRB seminar, February 2019, Collège de France, Paris, France.
- o Popgroup52, January 2019, Oxford, Royaume-Uni.

Poster

- Colloque annuel Society for Molecular Biology and Evolution (SMBE), July 2024, Puerto Vallarta, Mexico.
- o Population, Evolutionary, and Quantitative Genetics Conference, June 2022, Asilomar, USA.
- o Probabilistic Modelling in Genomics, October 2019, Aussois, France.
- o GDR AIEM, November 2018, Lille, France.
- o MCEB, June 2018, Saint-Martin-de-Londres, France.
- o GDR AIEM, November 2017, Lyon, France.

Media

2024 Interviews on the Indian Evolutionary History:

- General audience newspapers: Times of India
- Scientific newspapers: New scientist, Live Science
- o Radio/Podcast : Sher-E-Punjab AM 600 Vancouver

2021 Podcast episode of Chercheuses on my PhD journey: In French

Teaching Experience

- 2022 University of California, Berkeley, Quantitative Biology Bootcamp, Teaching Assistant (15h).
- 2022 University of California, Berkeley, Computational Biology Skill Seminar, lecture (1h, Video).
- 2018-2019 Sorbonne Université, Paris, Python, Teaching Assistant (40h).
- 2017-2018 IUT de Vélizy, C Language, class and tutorials (45h).

Mentoring/Supervision

2023-current Meaghan Marohn, PhD Candidate UC Berkeley.

Revisiting the evolution of lactase persistence: insights from South Asian genomes.

2024 Mara Baylis, Rotation student UC Berkeley, 3 months.

Description, origin and functional impact of the haplotype carrying the BTNL2 gene inherited from Denisovan and present in human populations.

2021 **Pierre Imbert**, M2 Bio-Informatique Paris Diderot, 6 months.

Benchmark of inference methods: a focus on population decline.

2019 **Ludovic Fourteau**, M1 Bio-Informatique Paris Diderot, 3 months.

Effect of a single recombination event on the diversity in a population.

2019 Rebecca Clodion, M1 Bio-Informatique Paris Diderot, 3 months.

The mysterious U spectrum: a population analysis of the genetic diversity.

Grants/Scholarship

2024 Symposium SMBE 2024 (Organisation grant 4,250\$).

2023 Keystone Symposia Future of Science Fund scholarship (Travel grant 1,200\$).

2020-2021 ATER Collège de France grant for a fourth year of PhD (Salary >20,000€).

2017-2020 Interfaces pour le vivant (IPV) doctoral school grant for a three year PhD (Salary >50,000€).

Service

Reviewer:

- American Journal of Human Genetics (2024)
- Molecular Biology and Evolution (2024)
- Human Population Genetics and Genomics (2024-2025)
- European Journal of Human Genetics (2022-2023)

Symposium organiser: *Haplotype-based methods and frameworks for inference of evolutionary history.* SMBE 2024, Puerto Vallarta, Mexico.

Student/postdoc representative of the ISYEB (elected member): Institute meetings participation. Organisation of monthly seminars and yearly Young Research day; from 2018 to 2021.

Jury: Best poster and best presentation of the symposium 'Evolutionary Ecology' of the 6th Young Natural History Scientists' Meeting, 12-16 mars 2019, MNHN, Paris, France.

Volunteer at conferences: Evolution 2018, Montpellier, France.

Student representative of the SDUEE Master (elected member): Council meetings participation; from 2016 to 2018.

Computer skills

Languages C, PYTHON

Familiar with the usage of a computer cluster

References/Collaborators

- Priya Moorjani, Postdoc advisor, Assitant Professor at UC Berkeley: moorjani@berkeley.edu
- o Guillaume Achaz, PhD advisor, Professor at *Université Paris-Cité*: guillaume.achaz@mnhn.fr
- Amaury Lambert, PhD advisor, Professor at École Normale Supérieure: amaury.lambert@ens.fr