Paul Bastide

Researcher in Statistics

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Current Position

2020 – present **Permanent CNRS Researcher**, *IMAG*, *Université de Montpellier*.

Keywords.

Phylogenetic Comparative Methods. Shift Detection. Model Selection. Phylogenetic Networks.

Past Position

2017 – 2020 **Postdoctoral researcher**, KU Leuven.

Bayesian Phylogenetic Comparative Methods applied to virology. With **Philippe Lemey** (KU Leuven) and **Marc Suchard** (UCLA).

Educational Background

2014 – 2017 **PhD Thesis**, *Université Paris Sud*.

Shifted stochastic processes evolving on trees: application to models of adaptive evolution on phylogenies. Supervised by **Stéphane Robin** (MIA, UMR 518 AgroParisTech/INRA) and **Mahendra Mariadassou** (MaIAGE, UR 1404 INRA).

2013 – 2014 Master 2 Math SV, Université Paris Sud - École Polytechnique - ENS Cachan.

Master in Mathematics For Life Sciences, awarded with highest honors.

2010 – 2014 Grande École, École Polytechnique.

(One of France's leading schools of science and engineering.) Majoring in Applied Mathematics.

Awards and Fellowships

2019 FWO long stay abroad grant, Research Foundation - Flanders.

Project based personal grant to visit Marc Suchard in UCLA.

2018 FWO Fellowship, Research Foundation - Flanders.

Three years project based personal post-doctoral fellowship.

2018 SFdS PhD Thesis Award, French Statistical Society.

Prix Marie-Jeanne Laurent-Duhamel, for a PhD in Applied Statistics, awarded every three years.

Fall 2015 Fulbright Scholar, University of Wisconsin - Madison.

To work with Cécile Ané at the Departments of Statistics and Botany, and in the Sytsma Lab.

Publications and Preprints

- 2020 S. Lequime, **P. Bastide**, S. Dellicour, P. Lemey, and G. Baele. nosoi: A stochastic agent-based transmission chain simulation framework in r. *Methods in Ecology and Evolution*, 11(8):1002–1007, jun 2020.
 - S. Dellicour, S. Lequime, B. Vrancken, M. S. Gill, **P. Bastide**, K. Gangavarapu, N. L. Matteson, Y. Tan, L. du Plessis, A. A. Fisher, M. I. Nelson, M. Gilbert, M. A. Suchard, K. G. Andersen, N. D. Grubaugh, O. G. Pybus, and P. Lemey. Epidemiological hypothesis testing using a phylogeographic and phylodynamic framework. *Nature Communications*, 11(1), nov 2020.

- **P. Bastide**, L. S. T. Ho, G. Baele, P. Lemey, and M. A. Suchard. Efficient Bayesian Inference of General Gaussian Models on Large Phylogenetic Trees. *Annals of Applied Statistics (in press)*, 2020, 2003.10336.
- G. Baele, M. S. Gill, **P. Bastide**, P. Lemey, and M. A. Suchard. Markov-modulated continuous-time markov chains to identify site- and branch-specific evolutionary variation in BEAST. *Systematic Biology*, may 2020.
- 2019 Z. Zhang, A. Nishimura, **P. Bastide**, X. Ji, R. P. Payne, P. Goulder, P. Lemey, and M. A. Suchard. Large-scale inference of correlation among mixed-type biological traits with phylogenetic multivariate probit models. *arXiv*, dec 2019, 1912.09185.
- 2018 **P. Bastide**, C. Solís-Lemus, R. Kriebel, K. W. Sparks, and C. Ané. Phylogenetic comparative methods on phylogenetic networks with reticulations. *Systematic Biology*, 67(5):800–820, apr 2018.
 - **P. Bastide**, C. Ané, S. Robin, and M. Mariadassou. Inference of Adaptive Shifts for Multivariate Correlated Traits. *Systematic Biology*, 67(4):662–680, jul 2018.
 - L. Aristide, **P. Bastide**, S. F. dos Reis, T. M. Pires dos Santos, R. T. Lopes, and S. I. Perez. Multiple factors behind early diversification of skull morphology in the continental radiation of New World monkeys. *Evolution*, 72(12):2697–2711, dec 2018.
- 2017 C. Solís-Lemus, **P. Bastide**, and C. Ané. PhyloNetworks: A Package for Phylogenetic Networks. *Molecular Biology and Evolution*, 34(12):3292–3298, dec 2017.
 - **P. Bastide**, M. Mariadassou, and S. Robin. Detection of adaptive shifts on phylogenies by using shifted stochastic processes on a tree. *Journal of the Royal Statistical Society: Series B (Statistical Methodology)*, 79(4):1067–1093, sep 2017, 1508.00225.
 - **P. Bastide**. *Shifted stochastic processes evolving on trees: application to models of adaptive evolution on phylogenies.* Phd thesis, Université Paris-Saclay, oct 2017.
- 2014 **P. Bastide** and T. David. Discours de réception d'Édouard Estaunié à l'Académie française : définitions croisées de la persona d'un académicien scientifique. *Épistémocritique*, 14(Greffes), 2014.

Softwares

- PhylogeneticEM (Main Author) An **R** package for automatic shift detection on phylogenies. Available on the CRAN and on GitHub.
- PhyloNetworks (Contributor) A **julia** package for statistical inference, data manipulation and visualization of phylogenetic networks. Available on GitHub.
 - BEAST (Contributor) A **java** cross-platform program for Bayesian analysis of molecular sequences and continuous traits using MCMC.
 - nosoi (Contributor) An **R** package for flexible agent-based stochastic transmission chain/epidemic simulation

Talks and Posters in Conferences

- 29/07/2019 **JSM 2019**, *Joint Statistical Meeting, Denver*, contributed speed talk.
- 05/06/2019 **JdS 2019**, 51èmes Journées de Statistique de la SFdS, Nancy, contributed talk.
- 21/08/2018 **Evolution 2018**, Joint Congress on Evolutionary Biology, Montpellier, contributed talk.
- 30/05/2018 **JdS 2018**, 50ème Journées de Statistiques de la SFdS, Saclay, **invited speaker** for the Prix Marie-Jeanne Laurent-Duhamel.
- 13/06/2017 MCEB 2017, Mathematical and Computation Evolutionary Biology, Porquerolles, poster.
- 30/05/2017 **JdS 2017**, 49èmes Journées de Statistique de la SFdS, Avignon, contributed talk.
- 28/06/2016 **JOBIM 2016**, Journées Ouvertes en Biologie, Informatique et Mathématiques, Lyon, contributed talk.

19/04/2016 **JPS 2016**, Jeunes Probabilistes et Statisticiens, École de Physique des Houches, talk. 22/06/2015 **MCEB 2015**, *Porquerolles*, poster. 02/06/2015 **JdS 2015**, 47èmes Journées de Statistique de la SFdS, Lille, contributed talk. Seminars and Workshops 12/10/2020 IMAG Statistics Seminar, Montpellier. 25/02/2020 LPSM Statistics seminar. Paris Diderot. 25/10/2019 TSU Statistics seminar, Texas State University. 25/04/2019 IMO Statistics seminar, Orsay. 21/01/2019 MaIAGE Statistics Seminar, Jouy-en-Josas. 19/10/2018 LBBE Statistics Seminar, Lyon. 28/09/2018 LaMME Statistics Seminar, Évry. 27/09/2018 LJK Probability & Statistics Seminar, Grenoble. 15/02/2018 Meeting of the group "Modélisation Mathématique et Biodiversité", Institut de Mathématique d'Orsay. 20/11/2017 Joint Seminar: "Mathématiques, Évolution, Biologie" and Statistics, Institut de Mathématiques de Marseilles. 08/11/2017 Bioinformatic Seminar (BIG), KU Leuven. 23/02/2017 Bioinformatics Team Meeting, INRA, Jouy-en-Josas. 14/11/2016 StatInfOmics Team Meeting, MaIAGE, Jouy-en-Josas. 04/10/2016 Journées des maths-info de l'INRA, Mallemort. 27/06/2016 AgroParisTech Statistics Seminar, Paris. 23/06/2016 Université Paris Sud Statistics Seminar (PhD Session), Orsay. 09/06/2016 Université Paris Descartes PhD Seminar, Paris. 18/05/2016 EDMH PhD Seminar, Paris. 11/04/2016 SupAgro Statistics Seminar, Montpellier. 10/02/2016 Sauquet Lab Botany Seminar, Université Paris Sud - Orsay. 19/11/2015 Sytsma Lab Botany Seminar, University of Wisconsin - Madison. 18/11/2015 University of Wisconsin Statistics Seminar, Madison. 22/07/2015 Morlon Lab Evolution Seminar, IBENS, Paris. 23/06/2015 SSB Statistics Seminar, Jouy-en-Josas. 16/02/2015 MaAIGE Internal Seminar, Jouy-en-Josas. **Teaching** 2018 **Teaching Assistant**, *KU Leuven*, in English and Dutch.

Applied Biostatistics. Master level, Biomedical sciences.

Introduction to Biostatistics. Bachelor level, Biomedical sciences.

2014 – 2017 **Teaching Assistant**, *Université Paris Sud*, in French.

Introduction to Mathematical Modeling. Bachelor level, Applied Mathematics (L3). [Springs 2015/16/17]

Introduction to Statistics. Bachelor level, Biology (L2). [Fall 2016]

Mathematics for Biology. Bachelor level, Biology (L1). [Fall 2016]

Measure Theory and Probabilities. Bachelor level, Mathematics (L3). [Spring 2016]

Introduction to Statistics. Bachelor level, Biology (L3). [Fall 2014]

2011 – 2012 **Tutor**, *GEPPM*.

A national program to help underprivileged high school students to continue their studies after high school. One course a week.

Implication in the Research Community

- Reviewer Systematic Biology, NeurIPS, PNAS, Evolutionary Bioinformatics, PeerJ, PlosOne, French ANR, Fulbright Program.
- Learned Society Member of the *Société Française de Statistique* (SFdS) and the *Society of Systematic Biologists* (SSB).
 - SFdS Member of the "comunication" focus work group of the Société Française de Statistiques.
 - Seminars Organizer of the IMAG Statistics seminar and the KIM Data and Life Sciences interdisciplinary seminar.
 - Outreach "Scoop" highlight on the network PCM paper for the Life Sciences Department of the Université Paris Saclay (www.scoop.it/t/life-sci-news-upsaclay?q=bastide)

Scientific Projects and Internships

- Spring 2014 Internship, UMR 518 AgroParisTech / INRA (Paris), Supervised by Stéphane Robin and Mahendra Mariadassou, Shifted Phylogenetic Comparative Methods.
- 2013 2014 **Project**, UMR de Génétique Végétale du Moulon (Saclay), Supervised by Christine Dillmann and Sylvie Huet, Analysis of the relations between the phenotype and the genotype on yeast organisms, combining genomic, proteomic and metabolic data.
- Spring 2013 **Internship**, International Institute for Climate and Society (IRI) at Columbia University (New York), Supervised by Rémi Cousin and Daniel Ruiz Carrascal, Construction and Study of a Multi-Model Framework For Short Term Predictions of a Malaria Epidemic in the Region of Kericho, Kenya.
 - Fall 2012 **Project**, CMAP (École Polytechnique), Supervised by Stéphane Gaubert, How to control the Chikungunya epidemic in La Réunion.
- 2011 2012 **Project**, CMAP (École Polytechnique), Supervised by Amandine Veber, Modeling the development of terrorists groups.

Other Work Experience

- Summer 2012 Internship, Intech-NSK, Novosibirsk (Russia).
 - 2010 2011 Internship, l'Enfant @ l'Hôpital, French association.

Skills

French First language

Programing R, Rcpp, Julia, Java, Matlab

English Spoken and written

Desktop tools MS Office, LaTex, Git

Notions Russian, Spanish, Dutch