Lorenzo Masoero

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Born July 20, 1992—Turin, Italy
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Education

2019-2021 PhD, Electrical Engineering and Computer Science, Massachusetts Institute of Technology
2017 - 2019 MSc in Electrical Engineering and Computer Science, Massachusetts Institute of Technology [1]
2015 - 2016 MA in Statistics and Applied Mathematics, with distinction, Collegio Carlo Alberto
2015 - 2016 MA in Quantitative Finance, 110/110 magna cum laude, Università degli Studi di Torino
2012 - 2014 BA in Economics, 110/110 cum laude, Università degli Studi di Torino

Scholarships and Awards

2020 SBSS Best Student Paper Award (ASA)
2020 Bayes Comp Travel Award
2018 BNP@NeurIPS Award
2017 Andrew (1956) and Erma Viterbi Fellowship
2016 Best Graduate Student of the Year (ATLEC)
2015 - 2016 Graduate Allievi Honors Program Scholarship, Collegio Carlo Alberto, Moncalieri
2012 - 2014 Undergraduate Allievi Honors Program Scholarship, Collegio Carlo Alberto, Moncalieri

Other Relevant Experience

2020 Applied Research Intern, Amazon CoreAI under the supervision of Professor Guido Imbens, Professor Thomas Richardson and Dr. James McQueen

Research

- "**Bayesian nonparametric strategies for power maximization in rare variants association studies**". **Spotlight** at Learning Meaningful Representations of Life, Neural Information Processing Systems, 2021 [poster]. Manuscript available on arXiv [https://arxiv.org/pdf/2112.02032.pdf] (M., Schraiber, Broderick)
- "**The Chicago Police Department dataset**". Neural Information Processing Systems Track on Datasets and Benchmarks, 2021, Dataset repository, current release (v0.1) and documentation (Horel, M., Agrawal, Roithmayr, Campbell)
- "**Scaled process priors: Improved predictions and uncertainties for new-feature counts via random scaling in Bayesian nonparametrics**". In submission. Featured in ISBA

[1] **Completed coursework**: Dynamic Programming and Stochastic Control (6.231) `[final project]`, Fundamentals of Probability (6.436), Inference and Information (6.437), Algorithms for Inference (6.438), Algorithmic aspects of Machine Learning (18.408) `[final project]`, Bayesian modeling and inference (6.882), Advanced stochastic processes (6.265), Mathematical Statistics: A Non-Asymptotic Approach (9.S914), Learning-Augmented Algorithms (6.890)

2021 [poster]. Manuscript available on arXiv [https://arxiv.org/pdf/2106.15480.pdf] (Camerlenghi, Favaro, M., Broderick)

- "**More for Less: Predicting and maximizing genetic variant discovery via Bayesian nonparametrics**"; to appear in Biometrika. **Best Student Paper Award**, awarded by the American Statistical Association, Section in Bayesian Statistical Science. Featured in SMEEB 2021, ASHG 2020, AABI 2019 [poster; presentation (YouTube)]; Manuscript available on arXiv [https://arxiv.org/pdf/1912.05516.pdf] (M., Camerlenghi, Favaro, Broderick)
- "**Independent finite approximations for Bayesian nonparametric inference: construction, error bounds, and practical implications**", in submission. Manuscript available on arXiv [https://arxiv.org/pdf/2009.10780.pdf] (Nguyen, Huggins, M., Mackey, Broderick)
- "**Posterior representations of hierarchical completely random measures in trait allocation models**", **Spotlight**, *BNP@NeurIPS2018* [poster] (M., Camerlenghi, Favaro and Broderick)
- "**Sensitivity of Bayesian inference to data perturbations**", *AABI 2018* [poster] (M., Stephenson, Broderick)
- "**Generic finite approximations for practical Bayesian nonparametrics**", **Spotlight**, *NIPS 2017 Workshop on Advances in Approximate Bayesian Inference* [poster]. (Huggins, M., Mackey, Broderick)

Theses

- "**Improved prediction and optimal sequencing strategies for genomic variant discovery via Bayesian nonparametrics**" — PhD thesis. Supervisor: Professor Tamara Broderick
- "**An asymptotic analysis of Gibbs-type priors**" — Master's thesis. Supervisors: Professors Pierpaolo de Blasi and Igor Prünster
- **"Econometrics of the Big Data"** — Undergraduate thesis. Supervisor: Professor Alessandro Sembenelli

Skills

- Proficient in Python (numpy, scipy, pandas, matplotlib, scikit-learn), LaTeX
- Past experience in C++, Matlab, R, RStudio

Talks, Poster sessions and Conference Presentations

2021

- CMS Statistics 2021, "Scaled process priors for Bayesian nonparametric estimation of the unseen genetic variation" [Invited session]
- ISBA: 2021 World Meeting of the International Society for Bayesian Analysis, "Scaled process priors for Bayesian nonparametric estimation of the unseen genetic variation" [Contributed session]
- SMEEB: Stochastic Models and Experiments in Ecology and Biology, "More for less: predicting and maximizing genomic diversity via Bayesian nonparametrics" [Contributed session]

2020

- American Society of Human Genetics meeting, "More for less: predicting and maximizing genomic diversity via Bayesian nonparametrics" [Poster session]
- Learning under complex structure, MIFODS workshop, *Cambridge (MA)*, "More for less: predicting and maximizing genomic diversity via Bayesian nonparametrics" [Poster session]
- Learning under complex structure, MIFODS workshop, *Cambridge (MA)*, "More for less: predicting and maximizing genomic diversity via Bayesian nonparametrics" [Poster session]
- Bayes Comp 2020, *Gainesville (FL)*, "More for less: predicting and maximizing genomic diversity via Bayesian nonparametrics" [Poster session]

2019

- Advances in Bayesian Nonparametric Methods and Its Applications, *Denver (CO), JSM 2019*, "Genomic variety prediction via Bayesian nonparametrics" [Topic-contributed session]
- Advances in Approximate Bayesian Inference, *Vancouver, Canada*, "More for less: Predicting and maximizing genetic variant discovery via Bayesian nonparametrics"
- Statistics and Data Science Conference 2019, *Cambridge (MA)*. "Genomic variety prediction via Bayesian nonparametrics"
- MLxMIT, *Cambridge (MA)*, "Genomic variety prediction via Bayesian nonparametrics"
- LIDS & Stats seminar, *Cambridge (MA)*, "Genomic variety prediction via Bayesian nonparametrics"
- CSAIL-MSR Trustworthy and Robust AI (TRAC) Workshop, *Cambridge (MA)*, "Getting the most bang for your buck: Predicting and maximizing the number of new genetic variants in a future experiment"

2018

- BNP@NeurIPS 2018, Montreal (Canada) "Posterior representations of hierarchical completely random measures in trait allocation models" [**Spotlight**]

Professional Service

2021 Reviewer for Statistical Science, AISTATS, ICBINB
2020 Reviewer for AAAI 2020, AISTATS 2020
2019 Reviewer for AISTATS 2019, NeurIPS 2019, AABI 2019
2018 Reviewer for BNP@NeurIPS2018