# Rajiv Boscolo Agostini, Ph.D.

https://raj94b.github.io/

## Research experience

11/2024 – ongoing Research Fellow. Life Sciences and Biotechnology Department, Population Genomics Research Group, University of Ferrara.

Statistical and bioinformatic analyses on ancient human genomic and metagenomic variability to reconstruct the Neolithic transition process in the Mediterranean basin.

12/2020 - 01/2021

**Collaborator.** Life Sciences and Biotechnology Department, Population Genomics Research Group University of Ferrara.

Application of machine learning algorithms on the analysis of genomic data of ancient and modern human populations.

04/2020 - 07/2020

**Research Fellow.** Life Sciences and Biotechnology Department, Population Genomics Research Group, University of Ferrara.

Hybrid genome assembly analyses of Tortoise (Testudo hermanni) data.

12/2019 - 10/2021

Attending Graduate. Life Sciences and Biotechnology Department, Population Genomics Research Group, University of Ferrara.

Genome assembly of the Appenine Brown Bear (Ursus arctos mariscanus) using long reads.

## **Education**

2021 – 2025 Ph.D., Evolutionary Biology and Ecology, University of Florence.

Curriculum: Genetics and Evolution

Thesis title: The evolutionary history of Italy from the Mesolithic to the present: lessons from

ancient genomes
Approved cum laude

2017 – 2019 M.Sc. Biomolecular and Evolutionary Sciences, University of Ferrara

Thesis title: Worldwide Helicobacter pylori genetic variation is informative about the evolu-

tionary history of anatomically modern humans.

Final mark: 110 cum laude

2014 – 2017 **B.Sc. Biological Sciences**, University of Ferrara

The sistifle: Worldwide genetic variability of Helicobacter pylori and identification of European

groups.

# Publications in peer-reviewed journals and preprints

A. Tommasi\*, **R. Boscolo Agostini**\*, G. Villani\*, N. Rambaldi Migliore\*, M. T. Vizzari, I. Caridnali, R. Di Gerlando, V. Nicolini, G. Sorasio, P. Santos, A. Olivieri, U. A. Perego, G. Catalano, N. Volante, L. Sarti, D. Caramelli, L. Sineo, H. Lancioni, A. Modi, S. Ghirotto\*\*, and A. Achilli\*\*, "Fifteen millennia of human mitogenome evolution in Sicily," (*Accepted*) Science Advances, 2025.

**R. Boscolo Agostini**, M. T. Vizzari, A. Benazzo, and S. Ghirotto, "Disentangling the worldwide invasion process of *Halyomorpha halys* through Approximate Bayesian Computation," *Heredity*, 2024. 
Ø DOI: 10.1038/s41437-024-00735-9.

F. F. Vale, **HpGP Research Network**, R. J. Roberts, I. Kobayashi, M. C. Camargo, and C. S. Rabkin, "Gene content, phage cycle regulation model and prophage inactivation disclosed by prophage genomics in the helicobacter pylori genome project," *Gut Microbes*, 2024. *O DOI:* 10.1080/19490976.2024.2379440.

- M. T. Vizzari, S. Ghirotto, **R. Boscolo Agostini**, P. Maisano Delser, L. Cassidy, A. Manica, and A. Benazzo, "Low-ABC: A robust demographic inference from low-coverage whole-genome data through abc," *BiorXiv*, 2024. ODI: 0.1101/2024.08.01.606209.
- K. Thorell, Z. Y. Muñoz-Ramírez, D. Wang, S. Sandoval-Motta, **R. Boscolo Agostini**, S. Ghirotto, R. C. Torres, D. Falush, M. C. Camargo, and C. S. Rabkin, "The *Helicobacter pylori* genome project: Insights into *H. pylori* population structure from analysis of a worldwide collection of complete genomes," *Nature Communications*, 2023. ODOI: 10.1038/s41467-023-43562-y.
- A. Modi\*, M. T. Vizzari\*, G. Catalano, **R. Boscolo Agostini**, S. Vai, M. Lari, C. Vergata, V. Z. Zaro, L. Liccioli, M. Fedi, S. Barone, L. Nigro, H. Lancioni, A. Achilli, L. Sineo, D. Caramelli, and S. Ghirotto, "Genetic structure and differentiation from early bronze age in the mediterranean island of Sicily: Insights from ancient mitochondrial genomes," *Frontiers in Genetics*, 2022. ODI: 10.3389/fgene.2022.945227.
- S. Sarno\*, **R. Boscolo Agostini**\*, S. De Fanti, G. Ferri, S. Ghirotto, G. Modenini, D. Pettener, and A. Boattini, "Y-chromosome variability and genetic history of commons from Northern Italy," *American Journal of Physical Anthropology*, 2021. ODOI: 10.1002/ajpa.24302.

## Contributions to conferences and meetings

- R. Boscolo Agostini, S. Vai, M. T. Vizzari, F. Santos Patricia De Pizzol, M. Vanni, A. Modi, A. Mittnik, E. Cilli, D. Luiselli, C. Martinez-Labarga, O. Rickards, N. Volante, L. Sarti, F. Martini, L. Sineo, D. Lo Vetro, D. Reich, S. Ghirotto, and D. Caramelli, *Unveiling the Complex Genetic History of Italy from Paleolithic to Middle Ages through Ancient Whole-Genome data*, Oral presentation, AAI-UZI Congress, Cagliari, 2025.
- **R. Boscolo Agostini**, La ricostruzione delle dinamiche popolazionistiche attraverso il DNA antico. Seminar, Department of Historical Sciences and Cultural Heritage, University of Siena, 2024.
- **R. Boscolo Agostini**, S. Vai, M. T. Vizzari, S. Perretti, F. De Pizzol, A. Mittnik, D. Luiselli, O. Rickards, L. Sineo, N. Volante, D. Reich, S. Ghirotto, and D. Caramelli, *The evolutionary history of Italy from Mesolithic to the Middle Ages: Lessons from ancient genomes*, Oral presentation, SIBE Congress 2024, University of Naples "Parthenope", 2024.
- **R. Boscolo Agostini**, S. Vai, M. T. Vizzari, S. Perretti, F. De Pizzol, A. Mittnik, D. Luiselli, O. Rickards, L. Sineo, N. Volante, D. Reich, S. Ghirotto, and D. Caramelli, *The evolutionary history of Italy from Mesolithic to the Middle Ages: Lessons from ancient genomes*, Poster Presentation, EMBO | EMBL Symposium, Heidelberg, 2024.
- S. Vai, **R. Boscolo Agostini**, A. Mittnik, D. Luiselli, O. Rickards, L. Sineo, N. Volante, D. Reich, S. Ghirotto, and D. Caramelli, 1000 Ancient Italian Genomes: Evidence from ancient biomolecules for unravelling past human population Dynamics, Tandem presentation, EAA Congress 2024, University of Rome La Sapienza, 2024.
- **R. Boscolo Agostini**, M. T. Vizzari, A. Benazzo, and S. Ghirotto, *Disentangling the worldwide invasion process of halyomorpha halys through ABC*, Poster presentation, SMBE, Ferrara, 2023.
- **R. Boscolo Agostini**, M. T. Vizzari, A. Benazzo, and S. Ghirotto, *Disentangling the worldwide invasion process of halyomorpha halys through ABC*, Joint Meeting AGI SIMAG 2023, Cortona, 2023.
- **R. Boscolo Agostini**, M. T. Vizzari, A. Benazzo, and S. Ghirotto, *Reconstructing the global invasion routes of Halyomorpha halys through Approximate Bayesian Computation*, Poster Presentation, SIBE Congress, Ancona 2022 and FISV, Napoli, 2022.

9

Y. Moodley, S. Ghirotto, A. Brunelli, **R. Boscolo Agostini**, N. De Serio, V. Chelysheva, A. Klyubin, A. Maady, K. Mominyalev, B. Linz, and M. Achtman, *The genetic history of modern humans in Siberia: A bacterial perspective*, Poster Presentation, SIBE Congress, Padova, 2019.

### **Outreach**

#### **Talks**

6/11/2023

**L'Ominazione: il giro del mondo in sei milioni di anni.** Talk presented in the outreach event "SCIENZE. PANORAMI SCIENTIFICI L'UOMO" hosted by Arci Rovigo

## Teaching and tutoring

#### **Academic tutor**

o2/2019 – o7/2019 Genetics. Genetics course in the Biological Sciences B.Sc. program, University of Ferrara. (55 hours)

11/2018 – 07/2019 **Biostatistics.** Tutoring for the Biostatistics course in the Biological Sciences B.Sc. program, University of Ferrara. (68 hours)

### Final thesis tutoring

2021 - 2023 Master thesis. Students: Giulia Battaglia, Sara Albertini.

2021 - 2022 **Bachelor thesis**. Students: Camilla Giacometti, Silvia Fiori, Brigida Pantaleo, Francesco Occhipinti.

# Workshps and courses

### Workshops

24/02/2021 - 26/02/2021

6/06/2023 - 9/06/2023 Phylogenetics workshop. Workshop organized by the Italian Society for Evolutionary Biology (SIBE) at the University of Trento.

12/06/2023 - 13/06/2023 The ancient DNA revolution: from the initial attempts to the Nobel Prize. Genetics school "Mario Polsinelli" organized by The Italian Geneticists Association (AGI) at Cortona.

31/07/2023 - 4/08/2023 Introduction to Ancient Metagenomics Summer School. Online workshop organized by SPAAM Community on data processing and ancient metagenomic analyses.

SIBE PHYLO Crash Course Online workshop by the Italian Society for Evolutionary Biology (SIBE), Exploring evolution using tree based methods: phylogenetics, molecular clocks, barcoding, and phylogenomics.

## **Skills**

Coding Bash, Python, R, Github

Genomic analyses Relatedness, Genomic structure, Phylogenomics, Coalescent simulations (fastsim-

coal, ms), Uniparental markers

Bioinformatics high-performance computing (HPC) clusters, reads processing, Genotype likeli-

hoods, Imputation, Variant calling, Genome assembly

Web Dev HTML, CSS,

Operating Systems | Windows, MacOS, Linux

Misc. Academic research, teaching, training, consultation.

### Other

### **Voluteering**

05/2023 - 07/2023 **Volunteer at SMBE 2023**.

Volunteer for the international scientific congress SMBE23 at Ferrara organized by the Society for Molecular Biology and Evolution (SMBE) and the University of Ferrara.

01/2025 - 09/2025 **Summer School Organizer**.

Organizer for the SIBE (Italian Society of Evolutionary Biology) Summer School 2025.

https://sites.google.com/view/sibesummerschool/home-page

Language certifications

2006 Certified Level B2 English (FCE). Awarded by Cambridge ESOL.

References

University of Ferrara (Italy) **Prof. Silvia Ghirotto**. email: silvia.ghirotto@unife.it

University of Ferrara (Italy) **Prof. Giorgio Bertorelle**. email: giorgio.bertorelle@unife.it