

Alba Refoyo-Martínez

PERSONAL DETAILS

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LinkedIn: <https://www.linkedin.com/in/albarefoyo>
Github: <https://github.com/albarema>

SKILLS

- Statistical data analysis
- Data management
- Data visualization
- Project management in cross-disciplinary groups
- Bioinformatics
- Software development
- Team leadership
- Mentoring

LANGUAGES

- English: proficiency
- Spanish: native
- Danish: beginner

EXPERTISE

- R
- R Markdown
- Python
- Jupyter Notebooks
- Conda
- Snakemake
- Bash scripting
- Git & Github

PERSONAL QUALITIES

- Motivated by challenges
- Proactive learner
- Highly organized
- Analytical
- Independent
- Decision-making
- Positive

SUMMARY

- 6 years' experience in bioinformatics, computational biology, and population genetics
- Demonstrated proficiency in the development of novel scientific software and reproducible open-source computational pipelines
- Deep expertise in scientific computing, using Python, R, snakemake and conda

EXPERIENCE

University of Copenhagen

Postdoctoral researcher

Nov 2022 - Present

- Developed novel computational tools and pipelines for processing genomic data
- Expanded the utility of R packages to specific research inquiries
- Conducted population genetic modelling in collaboration with Variant Bio

PhD student

Lundbeck Foundation Geogenetics Centre, Globe Institute

Oct 2019 - Oct 2022

Department of Integrative Biology at University of California, Berkeley

Sep 2022 - Oct 2022

- Computed statistical analyses and constructed computational pipelines for genomic data
- Programmed modelling of evolutionary complex processes
- Predicted genetic disease risk from whole-genome sequence data
- Implemented a tool to process tree-sequence data structures to track evolutionary events
- Performed data visualization of complex time-series genomic data

Master's student

Apr 2018 - Sep 2019

- Developed software in R to detect selection in complex demographic histories and evaluated its performance with state-of-the-art tools

EDUCATION

Doctor in Philosophy (DPhil)

Change of research environment

University of Copenhagen

2019-2022

University of California, Berkeley

Sep. 2022

Master of Science (MSc) in Bioinformatics

Elective student

University of Copenhagen

2017-2019

Copenhagen Business School

Bachelor of Science (BSc) in Health Biology

Erasmus program

Universidad de Alcalá

2013-2017

Vrije Universiteit Brussel

2016-2017

SCIENCE COMMUNICATION

I have contributed to numerous high-impact publications and developed my communication skills by presenting at global conferences.

- The Selection Landscape and Genetic Legacy of Ancient Eurasians. Irving-Pease, E.K.* , **Refoyo-Martínez, A.***, Ingason, A.* , [...], Racimo, F., Willerslev, E., 2022. *Nature accepted*
- Genetic risk for Multiple Sclerosis originated in Pastoralist Steppe populations. Barrie, W.* , Yang, Y.* , Atfield, K.E.* , Irving-Pease, E.K.* , E., Scorrano, G., Jensen, LT., Armen, A.P., Dimopoulos, EA., Stern, A., **Refoyo-Martínez, A.**, [...], Willerslev, E., 2022. *Nature accepted*
- Population Genomics of Stone Age Eurasia. Allentoft, M.E.* , Sikora, M.* , **Refoyo-Martínez, A.***, Irving-Pease, E.K.* , [...], Racimo, F., Kristiansen, K., Willerslev, E., 2022. *Nature accepted*
- How robust are cross-population signatures of polygenic adaptation in humans?. **Refoyo-Martínez, A.**, Liu, S., Jørgensen, A. M., Jin, X., Albrechtsen, A., Martin, A. R., Racimo, F., 2020. *Peer Community Journal*
- Identifying loci under positive selection in complex population histories. **Refoyo-Martínez, A.**, daFonseca, RR, Halldorsdottir, K, Arnason, E, Mailund, T, Racimo, F., 2019. *Genome Research*

VOLUNTEERING

- Conference chair (Evolutionary and Population Genetics in Denmark – EPIC 2023)
- Conference organizer (Evolutionary and Population Genetics in Denmark – EPIC 2022)
- Globe PhD Committee member organizing the GLOBE PhD Day 2022
- Course instructor in computational lab classes for PhDs in 'Analysis of large-scale datasets'
- Teaching assistant for bachelor students in 'Interaction design (KOMIT)' on the interaction and usability of user interfaces