

# Iker Rivas-González

PhD student at Aarhus University

PhD student in bioinformatics with experience in evolutionary and population genetics, data science and programming

EDUCATION		0
PhD in Bioinformatics Bioinformatics Research Centre, Aarhus University (Denmark) Supervised by Mikkel Heide Schierup	Feb 2020 – present	in
<ul> <li>Analyses of full genomes: primates, birds and marsupials.</li> <li>Phylogenetic reconstruction, selection, recombination.</li> <li>Mathematical modelling: hidden Markov models in population genetics.</li> <li>Data visualization.</li> </ul>		те
MSc in Bioinformatics Aarhus University (Denmark)	Aug 2018 – Jun 2021	
<ul> <li>Advanced genomics, big data analysis, data science, algorithmic engineering.</li> </ul>		
Erasmus exchange student Aarhus University (Denmark)	Aug 2017 - Jun 2018	•
<ul> <li>Erasmus scholarship, Basque Government scholarship, Kutxabank scholarship.</li> <li>Advanced molecular biology courses</li> </ul>		I۵
BSc in Biochemistry and Molecular Biology University of the Basque Country (Spain)	Sep 2014 - Jun 2018	
<ul> <li>With honors, extraordinary prize, best academic record of 2018.</li> <li>Protein engineering, biotechnology, physiology, metabolism.</li> </ul>		
WORK AND TEACHING EXPERIENCE		
Project: method development for modeling the evolutionary history of species	Nov 2021 – present	
<ul> <li>Developed a sequentially Markovian coalescent (SMC) to model the speciation process through incomplete lineage sorting (ILS).</li> </ul>		
Project: analysis of genome-wide data	Feb 2020 – present	
<ul> <li>Analyzed the ILS in the genomes of 50 primates, 250 birds and 6</li> </ul>		
<ul> <li>marsupials.</li> <li>Reconstructed the phylogeny of these species, successfully solving challenging nodes within the tree of life.</li> <li>Investigated the interplay between selection, recombination and ILS.</li> </ul>		
Software development in D	Nov 2020 - precept	
<ul> <li><u>PhaseTypeR</u>: phase-type distributions in R, available on CRAN.</li> <li><u>ggHoriPlot</u>: horizon plots in ggplot2, available on CRAN.</li> </ul>	Nov 2020 - present	
Teaching assistant, Aarhus University (Denmark)		
<ul> <li>Data Science in Bioinformatics (Fall of 2020 and 2021).</li> <li>Statistical and Machine Learning in Bioinformatics (Spring of 2020, 2021 and 2022).</li> </ul>		so
SELECTED PUBLICATIONS		-
<ul> <li>Iker Rivas-González, Marjolaine Rousselle, Fang Li, Long Zhou, Juli Munch, Yong Shao, Dongdong Wu, Mikkel H. Schierup and G Pervasive incomplete lineage sorting illuminates speciation and s</li> </ul>	en Y. Dutheil, Kasper Suojie Zhang (2021). election processes in	-

- Iker Rivas-González, Lars Nørvang Andersen and Asger Hobolth (2022). PhaseTypeR: phase-type distributions in R with reward transformations and a view towards population genetics. Journal of Statistical Software (under review).
- Shaohong Feng, Ming Bai, **Iker Rivas-González**, ..., Mikkel H. Schierup and Guojie Zhang (2022). Incomplete lineage sorting and phenotypic evolution in marsupials. *Cell;185(10):1646-1660.e18.* doi:<u>10.1016/j.cell.2022.03.034</u>.

## ATTENDED MEETINGS

primates. Science (under review).

- The Biology of Genomes (Cold Spring Harbor Laboratory, NY, USA) Talk
- Probabilistic Modelling in Genomics 2022 (Oxford, UK) Talk
- Three Minute Thesis Competition finalist (Aarhus, DK) Talk
- 6th Annual Danish Bioinformatics Conference 2021 (Aalborg, DK) Poster
- SMBE Meeting 2021 (virtual) Poster

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- github.com/rivasiker
- linkedin.com/in/rivasiker

#### CHNICAL SKILLS



## NGUAGE SKILLS

English Full professional proficiency

Full professional proficiency

Spanish

Native

Basque

Danish

Limited working proficiency 000German

> Elementary proficiency

## OFT SKILLS

- Leadership, teamwork, critical thinking
- Resilient, motivated, empathetic, cheerful
- Excellent written and verbal communication skills

### **INTERESTS**

- Graphic design
- Music production
- Vegetarian cuisine
- Basque folk dances
- Board games