# CURRICULUM VITAE March 2025

Osvaldo Antonio Martin

osvaldo.martin@aalto.fi https://orcid.org/0000-0001-7419-8978 https://aloctavodia.github.io/

# **Current Position**

#### **Research Fellow**

Aalto University. Finland. January 2025 – present. Conducting research with Prof. Dr. Vehtari focusing on advancing practical methods for Bayesian statistics and the Bayesian workflow.

## **Previous Positions**

### **Principal Data Scientist**

PyMC-Labs. February-December 2024. I helped companies solve challenging business problems by developing tailored Bayesian models and providing actionable insights through advanced data analysis.

#### Lecturer

Universidad Nacional de San Martín. Argentina. June-December 2024. Taught undergraduate students Bayesian statistics.

#### **Postdoctoral Researcher**

Aalto University. Finland. March 2021 – April 2022. Conducting research with Prof. Dr. Vehtari focusing on advancing practical methods for Bayesian statistics and the Bayesian workflow.

#### Researcher

National Scientific and Technical Research Council of Argentina (CONICET). Developed novel computational tools and methodologies in structural biology and Bayesian statistics. Mentored students in interdisciplinary research projects. Teach courses to undergrads and graduate students.

Investigador Independiente (Permanent position). April 2022 - June 2024.

Investigador Adjunto (Permanent position). 2017-2021

Investigador Asistente. 2014-2017

# **Academic Background**

#### PhD in Biology

National University of San Luis, Argentina 2007 – 2012.

Dissertation: *Chemical Shifts-Driven Monte Carlo Simulations*Focused on integrating computational methods with experimental data to determine and validate the structure of proteins and other biomolecules.

## **Publications** (last 5 years)

Full list at google scholar or ORCID. Google Scholar h-index 17

#### Recommendations for visual predictive checks in Bayesian workflow (PREPRINT)

Säilynoja T, Johnson A.R. Martin, O.A. Vehtari A. ArXiv 2025.

### Prior knowledge elicitation: The past, present, and future

Mikkola, P. Martin, O.A. Chandramoul, S. Hartmann, M. Abril-Pla, O. Thomas, O. Pesonen. Corander, J. Vehtari, Aki. Kaski S. Bürkner, PC. Klami A. Bayesian Analysis. 2024.

#### PvMC: A modern, and comprehensive probabilistic programming framework in Pvthon

Abril-Pla O, Andreani V, Carroll C, Dong L, Fonnesbeck CJ, Kochurov M, Kumar R, Lao J, Luhmann CC, Martin O.A., Osthege M, Vieira R, Wiecki T, Zinkov R.. PeerJ Computer Science 9:e1516. 2023.

## PreliZ: A tool-box for prior elicitation

Icazatti A, Abril-Pla O, Klami A. Martin O.A.. The Journal of Open Source Software. 2023.

# Modeling Rainwater Harvesting and Storage Dynamics of Rural Impoundments in Dry Chaco Rangelands

Niborski M.J. Martin, O.A. Murray F. Jobbágy EG Nosetto MD Paez RA Magliano PN. Water. 2023.

#### Bambi: A simple interface for fitting Bayesian linear models in Python

Capretto, T. Piho, C. Kumar, R. Westfall, J. Yarkoni, Tal. <u>Martin O.A.</u> Journal of Statistical Software. 2022.

# A call for changing data analysis practices: from philosophy and comprehensive reporting to modeling approaches and back

Martin O.A, Teste F.P. Plant and Soil. 2022.

## Exploring the quality of protein structural models from a Bayesian perspective

Arroyuelo, A. Vila, J.A. Martin O.A. Journal of Computational Chemistry. 2021.

### Bayesian additive regression trees for probabilistic programming (PREPRINT)

Quiroga M Garay P. G. Alonso J. M., Loyola J. M. Martin O.A. ArXiv 2022.

#### Books

### **Bayesian Analysis with Python**

Martin O.A. Packt Publishing. Third Edition. ISBN-13: 978-1805127161. 2024.

# **Bayesian Modeling and Computation in Python.**

Martin O.A. Kumar R, Lao J. CRC press. ISBN 978-0367894368. 2021.

# Online teaching material

#### **Practical MCMC**

Intuitive Bayes. Online learning platform for Bayesian statistics. (paid)

#### **Bayesian modelling**

https://gmp.net.ar/modelado bayesiano (free, Spanish)

# **Exploratory Analysis of Bayesian Models**

https://arviz-devs.github.io/Exploratory-Analysis-of-Bayesian-Models (free, work in progress)

# Past teaching experience

#### Lecturer

Bayesian Modelling. National University of San Luis. Hourly credit 60 hours. 2022, 2023

**Bayesian data analysis**. National University of Tucumán. Second week March 2020

Bayesian data analysis. National University of San Luis. First week December 2019

Advanced methods in statistics and data analysis. National University of San Luis October-December 2019

**Introduction to data science**. National University of San Luis. Hourly credit 60 hours. February-March 2018.

**Bayesian data analysis**. National University of San Luis Hourly credit 60 hours. August-September 2018.

#### **Teaching Assistant**

Bavesian Data Analysis. Aalto University. 2021

Structural Bioinformatics. UNSL. Hourly credit 90 hours. 2013-2016.

Algorithms in bioinformatics. UNSL. Hourly credit 80 hours. February 2015.

# Funding as main researcher/principal investigator

Bayesian additive regression trees: automatic inference, variable selection, and interpretability. PICT-GRF (2023-2024). Note: This is a competitive research grant in Argentina (~\$12,000 USD), the highest available for my career level within the national funding system.

**Dashboard-assisted exploratory analysis of Bayesian models**. NumFOCUS small development grants. (2023). \$10,000 USD

Jax SMC samplers for PyMC. NumFOCUS small development grants. (2022). \$10,000 USD

Fine tuning the Bayesian Additive Regression Trees implementation in PyMC. NumFOCUS small development grants. (2021). \$10,000 USD

Improving and expanding the implementation of Approximate Bayesian Computation. NumFOCUS small development grants. (2020). \$10,000 USD

Create educational material and give workshops related to exploratory analysis of Bayesian models with ArviZ. NumFOCUS small development grants. (2019). \$10,000 USD

**Probabilistic programming for Structural Bioinformatics**. PICT-Joven (2017-2019). Note: This is a competitive research grant in Argentina (~\$5,000 USD), the highest available for my career level within the national funding system.

#### Advisor and co-advisor

#### **Doctoral students**

Co-Advisor: Noa Kallioinen. *Tools for choosing and analysing priors in Bayesian modelling*. Aalto University. (2021-). Ongoing.

Advisor: Agustina Arroyuelo. *Structural determination of biomolecules by statistical inference*. National University of San Luis. (2017-2021). Approved.

Co-advisor: Alejandro Icazatti. *Validation and determination of the structure of nucleic acids*. National University of San Luis. (2017-2021). Approved

Co-advisor: Pablo Garay. Study of carbohydrate conformational preferences using chemical shifts. National University of San Luis. (2013-2017). Approved

#### **Bachelor students**

Advisor BSc Thesis: Franco Joaquin Loyola. *Application of optimization methodologies on Bayesian Additive Regression Trees*. Computer engineering. National University of San Luis. Ongoing.

Advisor BSc Thesis: Agustina Arroyuelo. *Development of a software application for creating, viewing and analyzing models of macromolecules*. Molecular Biology. National University of San Luis. 2016. Approved.

Advisor BSc Thesis: Pedro Ramírez. *Protein structure refinement guided by* <sup>13</sup>C<sup>a</sup> y <sup>13</sup>C<sup>b</sup> chemical shifts. Molecular biology. National University of San Luis. Approved December 2014.

Co-advisor BSc Thesis: Alejandro Icazatti. *Combined use of chemical shifts of 13Cα and 13Cβ in protein validation.* Molecular Biology. National University of San Luis. Approved 2013.

Co-advisor BSc Thesis: Ezequiel Frigini. *Is the cell membrane, permeable to the diffusion of glyphosate?* Molecular biology. National University of San Luis. Approved December 2014.

# Software development

#### Core developer

ArviZ. Exploratory analysis of Bayesian models

Bambi. Bayesian Model-Building Interface in Python.

<u>Kulprit</u>. Kullback-Leibler projections for Bayesian model selection

PreliZ. A tool-box for prior elicitation

PyMC. Probabilistic Programming in Python

PyMC-BART. Bayesian Additive Regression Trees for Probabilistic programming

# Older projects

*CheShift*: A web-server for protein-structure validation.

**PyMOL** Plugins: PyMOL is a molecular visualization system

# **Synergistic activities**

Member of the PhD in Biology Committee. National University of San Luis. 2019-2021.

Head of the organizing committee of Argentinean workshop of scientific computation, San Luis, Argentina 2019.

Head of the organizing committee of PyData, San Luis, Argentina 2017. This was the first PyData in Latin-America. Member of the PyData San Luis chapter.

Co-host La Búsqueda, popular science radio-show and podcast. Radio UNSL 2008-2009. Radio Ciudad 2010