CURRICULUM VITAE

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Academic Background

PhD in Biology. *Chemical Shifts-Driven Monte Carlo Simulations*. National University of San Luis. 2007-2012.

Research positions

Postdoc Researcher. Aalto University. March 2021 – April 2022 *Investigador Adjunto*. National Scientific and Technical Research Council (CONICET). 2017-present *Investigador Asistente*. National Scientific and Technical Research Council (CONICET). 2014-2017

Recent scientific publications

Only last five years, visit google scholar for the entire list.

- Capretto, T. Piho, C. Kumar, R. Westfall, J. Yarkoni, Tal. **Martin O.A.** *Bambi: A simple interface for fitting Bayesian linear models in Python*. Journal of Statistical Software (2022).
- **Martin O.A**, ; Teste F.P. A call for changing data analysis practices: from philosophy and comprehensive reporting to modeling approaches and back. Plant and Soil (2022).
- Arroyuelo, A. Vila, J.A. **Martin O.A.** *Exploring the quality of protein structural models from a Bayesian perspective.* Journal of Computational Chemistry (2021).
- **Martin O.A**, Vila J.A. *The Marginal Stability of Proteins: How Jiggling and Wiggling of Atoms are Connected to Neutral Evolution*. Journal of molecular evolution (2020).
- Arroyuelo A., **Martin, O.A** Scheraga H.A. Vila, J. A. *Assessing the One-Bond C\alpha-H Spin-Spin Coupling Constants in Proteins: Pros and Cons of Different Approaches*. The Journal of Physical Chemistry B. (2020)
- Icazatti A.A, Loyola J.M., Szleifer I, Vila J.A. **Martin O.A.** *Classification of RNA backbone conformations into rotamers using 13C' chemical shifts: exploring how far we can go.* PeerJ 7, e7904 (2019)
- **Martin O.A.** Vorobjev Y, Scheraga HA, Vila JA. Outline of an experimental design aimed to detect protein A mirror image in solution. PeerJ Physical Chemistry 1, e2 (2019)
- Baldi G, Schauman S.A, Texeira M, Marinaro S, **Martin O.A**, Gandini P, Jobbágy E.G. *Nature representation in South American protected areas: country contrasts and conservation priorities.* PeerJ 7, e7155 (2019)
- Kumar, R. Carroll C, Hartikainen A, **Martin, O.A** *ArviZ a unified library for exploratory analysis of Bayesian models in Python*. The Journal of Open Source Software (2019)
- Icazatti, A. **Martin, O.A**. Villegas, M. Szleifer, I. Vila, J.A.. *13Check_RNA: A tool to evaluate 13 C chemical shifts assignments of RNA*. Bioinformatics (2018)
- Alonso, J.M. Arroyuelo, A. Garay, P.G. **Martin, O.A**. Vila J.A. *Finite Dimension: A Mathematical Tool to Analise Glycans*. Scientific Reports (2018)
- Garay, P.G. Vila, J. A. **Martin, O.A**. *CheSweet: An application to predict glycan's chemicals shifts*. The Journal of Open Source Software (2018)

Preprints

- 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. Prior knowledge elicitation: The past, present, and future arXiv (2021)
- Quiroga M, Garay P. G. ,Alonso J. M. , Loyola J. M., and **Martin O.A.** *Bayesian additive regression trees for probabilistic programming.* arXiv 2022.

Books

Bayesian Modeling and Computation in Python. Martin O.A. Kumar R, Lao J. CRC press. ISBN 978-0367894368. 2021.

Bayesian Analysis with Python. Martin O.A. Packt Publishing. Second Edition. ISBN-13: 978-1789341652. 2018

Research projects funding as main researcher/principal investigator

Create educational material and give workshops related to exploratory analysis of Bayesian models with ArviZ. NumFOCUS small grants. (2019)

Probabilistic programming for Structural Bioinformatics FONCYT. PICT-Joven (2017-2019)

Advisor for undergrad and graduate students

Current projects

PhD Advisor: Lic. Miriana Quiroga. *Bayesian additive regression trees: automatic inference, variable selection and interpretability*. National University of San Luis. (2020-2024)

Previous projects

PhD Advisor: Lic Agustina Arroyuelo. *Structural determination of biomolecules by statistical inference*. National University of San Luis. (2016-2021)

Co-advisor Phd Thesis: Alejandro Icazatti. *Validation and determination of the structure of nucleic acids*. National University of San Luis. (2014-2019).

Co-advisor Phd Thesis: Pablo Garay. *CheSweet: Determination and validation of glycans and glycoproteins*. National University of San Luis. (2013-2017).

Advisor BSc Thesis: Agustina Arroyuelo. *Development of a software application for creating, viewing and analyzing models of macromolecules*. Thesis to get the degree of Bachelor in Molecular Biology. National University of San Luis. Approved September 2019.

Advisor BSc Thesis: Pedro Ramírez. *Protein structure refinement guided by* ¹³*C*^a *y* ¹³*C*^b *chemical shifts*. Thesis to get the degree of Bachelor in Molecular biology. National University of San Luis. Approved December 2014.

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

Software development

Current projects

Core developer of ArviZ. Exploratory analysis of Bayesian models

Core developer of **Bambi**. Bayesian Model-Building Interface in Python.

Core developer of **PyMC**. Probabilistic Programming in Python

Older projects

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

<u>PyMOL</u> 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 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

