Verónica Miró Pina

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Born in Murcia (Spain), February 18th 1991 ⊠ veronica.miro@crg.eu

	Current position
2020	
2020	Postdoc , Weghorn's lab, Centre for Genomic Regulation, Barcelona (Spain).
	Research Interests
	Cancer genomics, Population genetics, Evolutionary biology.
	Deep Learning, Neural Networks.
	Statistical inference, Data-analysis, Simulations.
	Probability theory, Stochastic processes, Coalescent theory.
	Previous positions
2018 - 2020	Postdoc , Instituto de Investigaciones en Matemáticas Avanzadas y Sistemas (IIMAS), Universidad Nacional Autónoma de México (UNAM).
	Education
2015 - 2018	PhD , Equilibrium patterns of genetic diversity shuffled by migration and recombination, LPSM (Sorbonne Université) and CIRB (Collège de France), Advisors : Emmanuel Schertzer and Amaury Lambert.
2013-2014	Masters (2nd year), Applied Mathematics, Université Pierre et Marie Curie, Paris 6, with honors.
2012-2013	Master (1st year), Biology, École Normale supérieure (Paris).
2011-2012	Bachelors Degree, Biology, École Normale supérieure (Paris).
	Publications
	Published papers
2022	Segregational instability of multicopy plasmids: A population genetics approach,

- Hernandez, J.C.R.*, Miró Pina, V.*, Siri-Jégousse, A., Peña Miller, R., Palau, S. and González Casanova, A., PLoS ONE, 12(12), e9469.
 * both should be considered first authors
- 2022 The role of connectivity on COVID-19 preventive approaches, Miró Pina*, V,. Nava-Trejo*, J. Tobiás, A., Nzabarushimana, E., González Casanova, A. and González Casanova, I., PLoS ONE, 17(9): e0273906.
 * both should be considered first authors
- 2022 The symmetric coalescent and Wright-Fisher models with bottlenecks, González Casanova, A., Miró Pina, V. and Siri-Jégousse, A., Annals of Applied Probability, 32(1), 235– 268.

† authors listed in alphabetical order

- 2022 Estimating the time since admixture from phased and unphased molecular data, Janzen, T., Miró Pina, V., Molecular Ecology Resources, 22(3), 908–926.
- 2021 Chromosome Painting, Lambert, A., Miró Pina, V. and Schertzer, E., Annals of Applied Probability, 31(2), 826–864.
 † authors listed in alphabetical order
- 2020 The Wright-Fisher model with efficiency, González Casanova, A., Miró Pina, V. and Pardo, J.C., Theoretical Population Biology, 132, 33–46.
- 2019 How does geographic distance translates into genetic distance?, Miró Pina, V and Schertzer, E., Stochastic processes and their Applications, 129(10) 3839-3921 7.
 † authors listed in alphabetical order

Preprints

- 2023 An extension of the Walsh-Hadamard transform to calculate and model epistasis in genetic landscapes of arbitrary shape and complexity, Faure, A.J., Lehner, B., Miró Pina, V., Serrano Colomé, C. and Weghorn, D., bioRxiv 2023.03.06.531391, under revision. † authors listed in alphabetical order
- 2022 The stochastic speed of coming down from infinity for general Dirichlet Xi-coalescents, González Casanova, A., Miró Pina, V., Schertzer, E. and Siri-Jégousse, A., aRxiv 2209.13438, under revision.
 † authors listed in alphabetical order

Grants and fellowships

- 2022-2023 Agaur LLavor (2021 LLAV 00061), Project grant for innovative projects with the potential for being incorporated into the production sector, from the Catalan Agency for Management of University and Research Grants, project: TAFI (Tumor Allele Frequency Interpreter) : a Deep Learning tool to reveal tumor heterogeneity.
- 2018-2022 Postdoctoral fellowship, DGAPA-UNAM (Universidad Nacional Autónoma de México).
- 2015-2018 PhD scholarship, PhD scholarship for students from École Normale Supérieure.

Research Experience

- May-July Laboratoire de Biométrie et Biologie Évolutive, University Lyon 1 (Lyon, France),
 2015 Nicolas Lartillot, Study of an individual-based model of species diversification.
 Statistical inference, MCMC algorithms
- January- Department of Ecology and Evolutionary Biology, Princeton University, Simon April Levin's Lab, Theoretical ecology, Study of the speciation mechanisms. 2015
- February SMILE team (Stochastic Models for the Inference of Life and Evolution), CIRB,
- 2014 Collège de France, Emmanuel Scherter and Amaury Lambert, Study of a population-based July 2015 model of speciation.
- February Center for Genomics and Systems Biology, NYU, New York University, Edo Kussell
- -June 2013 Experimental evolution.
- June-July Centro Nacional de Biotecnología, CSIC, Universidad Autónoma de Madrid (Spain), 2012 Juan Poyatos

Experimental study of a genetic network in E.coli.

Teaching experience

- 2022 Supervisor. Larisa Arreola's undergraduate tesis, UNAM.
- 2020 Supervisor. Fernanda López Eslava's undergraduate, UNAM.
- 2020 Supervisor. Amilkar Gazque's undergraduate student, UNAM.
- 2019 Teacher. Probability 2 and Applied mathematics seminar. Undergrad level at UNAM
- 2015-2018 Teaching assistant at Ecole Normale Supérieure (Paris): Mathematics for biologist. Master's level
- 2015-2018 Teaching assistant at Université Pierre et Marie Curie (Paris): Probability, Differential equations, Analysis. Undergrad level.
- 2012-2015 Oral examiner at lycée Henri IV (Paris, France). Undergrad level

Selected Presentations

- 2022 TAFI (Tumor Allele Frequency Interpreter): a new deep learning tool to reveal the evolutionary history of tumors. Contributed talk, Probgen 2022, Oxford
- 2022 TAFI (Tumor Allele Frequency Interpreter): a new deep learning tool to reveal the evolutionary history of tumors. Poster, WE-Heraeus-Seminar / Evolution of Cancer Reconstructing the Past, Predicting the Future
- 2021 Estimating the time since admixture from phased and unphased molecular data. Poster SMBE 2021 Society for Molecular Biology and Evolution
- 2021 Ξ-coalescents arising from population models with bottlenecks. Journées MAS, Institut Denis Poisson, Orléans, France.
- 2021 Ξ-coalescents arising from population models with bottlenecks. Probability seminar, University of Bath, UK (online).
- 2020 The symmetric coalescent. Bernoulli-IMS One World Symposium 2020. Available on YouTube.
- 2019 The symmetric coalescent. Invited talk. CLAPEM 2019, Merida, Yucatan, Mexico

- 2019 A population genetics model to understand plasmid loss and maintenance, segundo taller nacional de Probabilidad y Biologia, Cuernavaca
- 2019 The symmetric coalescent. Invited talk. SPA 2019, Chicago, US
- 2019 The symmetric coalescent. MMEE 2019, Lyon, France
- 2019 The symmetric coalescent. Invited talk. Congreso de la Sociedad Matemática Mexicana, Monterrey, México.
- 2019 Spatial models in Population Genetics. Workshop Ecosystem dynamics: stakes, data and models, Institut Pascal, Saclay, France
- 2018 Chromosome Painting. Primer taller nacional de Probabilidad y Biologia, Cuernavaca.
- 2018 Chromosome Painting. Saint Flour Probability School France).
- 2017 Chromosome Painting. AIEM SMBE meeting (regional meeting of the Society for Molecular Biology and Evolution). Lyon (France).
- 2017 How does geographic distance translates into genetic distance? Annual meeting of the French research group 'Theory and Models of Biodiversity' Paris (France).
- 2017 Chromosome Painting. 'Les Probabilités de demain' (young researchers in Probability meeting), IHES (France).
- 2017 How does geographic distance translates into genetic distance? Conference MMEE (Mathematical Models in Ecology and Evolution), London (UK).
- 2016 Chromosome Painting (poster) Conference MCEB (Mathematical and Computational Evolutionary Biology), Montpellier (France).
- 2016 How does geographic distance translates into genetic distance? ECMTB (European Conference on Mathematical and Theoretical Biology), Nottingham (UK).

Other skills

Computer Python, R, bash (advanced), C++, Matlab (basics) science

Languages Bilingual in Spanish and French. Fluent in English. Basic knowledge of Catalán

Organisation 2021-2022. Postdoc representative: member of the CRG postdoc comittee, organisation of carreer related and social events.

2015-2016. Student representative: organisation of cultural and social events for the students and postdocs at Collège de France.

2011-2012. Student representative: organisation of social events and of the annual gala at Ecole Normale Supérieure.