

# Irene Marín Goñi

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## Education

### Mayo Clinic

Visiting Graduate Student

📅 Nov 2022 - Apr 2025 📍 USA

- **Cancer and translational Certificate courses:** Genomic Analysis of Complex Traits, Molecular Variant Evaluation, Cancer Biology, Molecular Genetics, Drug Metabolism, Molecular Pharmacology of Receptor Signaling.
- **Advisor:** Liewei Wang MD, PhD (*Pharmacogenomics and Drug Targets Lab, Mayo Clinic*)

### University of Navarra

PhD in Computational Biology

📅 Sep 2021 - Currently 📍 Spain

- **Research topic:** Understanding molecular mechanisms of abiraterone resistance in metastatic castration resistant prostate cancer.
- **Courses:** Bioinformatics, Statistics, Research Ethics, Advanced Computational Biology methods, Experimental Animal Work, Intellectual Property.
- **Teaching:** Biostatistics, Introduction to Chemistry and Nutrition.
- **Advisors:** Mikel Hernaez PhD (*Machine Learning in Biomedicine Lab, CIMA*) and Liewei Wang MD, PhD

MSc in Computational Methods in Science

📅 Sep 2020 - Sep 2021 📍 Spain

- **Courses:** Computer Programming Languages, Biostatistics, NGS data processing and analysis, Machine Learning and Deep Learning (Intro), Data Analytics.

Joint program Master's of Pharmacy and BSc in Human Nutrition and Dietetics

📅 Sep 2013 - Feb 2020 📍 Spain

- **Pharmacy Internships:** 3 months at community pharmacy Greenlight Pharmacy-Stepney (UK), 3 months at hospital pharmacy Clinica Universidad de Navarra (Spain) and 1 month in Hospital Clinical Analysis Laboratory (Spain).
- **International Exchange Programs:** Cornell University (USA), University of Ipswich (UK), Amsterdam University of Applied Sciences (NL).
- **GPA:** 3.45 or 9.32/10 (**Top 5** in Pharmacy) and 3.4 or 9.23/10 (**Top 1** in Nutrition)
- Senior Class Student Representative.

## Research Experience

### Wet lab Research projects

PhD thesis project at Mayo Clinic

📅 Sep 2022 - Dec 2024 📍 Spain

- **Main project:** Targeting of CDK12/13 and androgen pathways in metastatic castration-resistant prostate cancer. **PI:** Dr.Wang.

Research Summer Intern at Navarrabiomed

📅 Jun 2021 - July 2021 📍 Spain

- **Cancer Signaling Lab:** Study of the effects on the expression of PD-L1 in glioma and breast cancer cells of ranolazine, a disruptor of the lipid metabolism. **PI:** Imanol Arozarena PhD

Undergraduate research projects at University of Navarra

📅 2016 - 2020 📍 Spain

- **PharmD Degree Thesis:** Evaluation of the cytotoxic activity of solid lipid nanoparticles loaded with edelfosine in a neuroblastoma model.
- **BSc Nutrition Degree Thesis:** Evaluation of the nutritional regulation of PRDM1 in murine adipocytes.
- **Research Intern:** Evaluation of organoselenium compounds in lipolysis proteins in murine adipocytes.

### Computational Research projects

Human Tumor Atlas Network (HTAN)

📅 November 6-8, 2024 📍 NIH Campus, Bethesda, MD

- Attended in-person to the **HTAN Data Jamboree**. The goal is to collaboratively build unique solutions that solve relevant problems in cancer research using NCI CRDC cloud resources and HTAN data and public datasets (**spatial omics and single cell**). *Manuscript in progress*

Mayo Clinic collaborations

📅 Nov 2022 - Current 📍 Rochester, MN (USA)

- **High throughput drug screening in PDX models** of metastatic castration resistant prostate cancer patients to overcome abiraterone resistance. **PI:** Liewei Wang MD, PhD.
- **Analysis of RNAseq data** of LSD1 inhibitors for prostate cancer treatment **PI:** Dr.Wang.
- **Analysis of single nuclei-RNAseq data** of syngeneic model of triple negative breast cancer treated with DNMTi and immunotherapy. **PI:** Dr.Wang.
- **Spatially resolved transcriptomics** of cardiac fibrosis associated with dilated cardiomyopathy in transgenic mice. **PIs:** Naveen Pereira, MD. and Richard Weinshilboum MD, PhD.
- **Other collaborations:** Bioinformatic support for alcohol use disorder project. **PIs:** Duan Liu, PhD. and Richard Weinshilboum MD, PhD.

- **Main PhD research topic:** Understanding molecular mechanisms of abiraterone resistance in metastatic castration resistant prostate cancer through bulk, single cell and spatial transcriptomics.
- **MSc Dissertation project:** Study of the differential gene regulatory networks in the context of metastatic castration-resistant prostate cancer and abiraterone treatment. Collaboration with Mayo Clinic.
- **Other collaborations:**
  - Interpretable Causal Representation Learning for Biological Data in the Pathway Space
  - Sweetwater: An interpretable and adaptive autoencoder for efficient tissue deconvolution
  - Study of gene regulatory networks with Quantum Computing.

## Research Scholarships

### Postgraduate program fellowship

LaCaixa Foundation

 Aug 2023 - Dec 2024
  USA

2-year excellence fellowship for studies abroad. Intended to support my research stay at Mayo Clinic.

### PhD studies fellowship

Spanish Government

 Dec 2023 - Currently
  Spain

4-year full-time Government-funded scholarship for PhD student stipend.

### PhD studies fellowship and mobility fellowship



Government of Navarra

 Feb 2023 - Dec 2023
  Spain

3-year full-time funded scholarship for PhD student tuition and stipend with additional support for research stay in Mayo Clinic.

### PhD studies fellowship

CIMA University of Navarra

 Sep 2022 - Feb 2023
  Spain

Funded through Congressionally-directed medical research program (CDMRP) grant (US Department of Defense)

## Honors and Awards

2020	<b>Special End of Studies Award</b> , for BSc in Human Nutrition and Dietetics, recognition for <b>Top 1</b> student of each major class (GPA: 3.4 or 9.23/10)	University of Navarra
Fall 2018	<b>Dean's List</b> , College of Human Ecology (GPA 4.162)	Cornell University
2017 - 2018	<b>Senior Class Student Representative</b> , elected by Pharmacy School classmates	University of Navarra

## Publications

### 2025

Interpretable Causal Representation Learning for Biological Data in the Pathway Space

 Jesús de la Fuente, Robert Lehmann, Carlos Ruiz-Arenas, **Irene Marín-Goñi**, Jan Voges, Xabier Martinez de Morentin, David Gomez-Cabrero, Idoia Ochoa, Jasper Tegnér, Vincenzo Lagani, Mikel Hernáez
Accepted manuscript (*ICLR*) (2025)

FNDC4 Alters Glutamatergic and GABAergic Neurogenesis

 Xiujuan Zhu, John J August, Li Wang, Kim Soan, Enci Ding, Ateka Saleh, **Irene Marín-Goñi**, Huanyao Gao, Ching Man Wai, Irene Moon, Brandon Coombes, Tony M Kerr, Noboyushi Suto, Liewei Wang, Mark A. Frye, Joanna Biernacka, Victor Karpyak, Hu Li, Richard M. Weinshilboum, Duan Liu
Under revision in *Journal of Clinical Investigation* (2025)

Sweetwater: An interpretable and adaptive autoencoder for efficient tissue deconvolution

 Jesús de la Fuente, Naroa Legarra, Guillermo Serrano, **Irene Marín-Goñi**, Aintzane Diaz-Mazkaran, Markel Benito Sendin, Ana Garcia Osta, Krishna R. Kalari, Carlos Fernandez-Granda, Idoia Ochoa, Mikel Hernáez
arXiv. Under second review at *Nucleic Acid Research*, 2023 (2024)

### 2024

NetActivity enhances transcriptional signals by combining gene expression into robust gene set activity scores through interpretable autoencoders

Carlos Ruiz-Arenas, **Irene Marín-Goñi**, Liewei Wang, Idoia Ochoa, Luis A Pérez-Jurado, Mikel Hernaez*Nucleic Acids Research* 52.9 (2024) e44–e44

Bayesian machine learning enables identification of transcriptional network disruptions associated with drug-resistant prostate cancer  
Charles Blatti, Jesús de la Fuente, Huanyao Gao, **Irene Marín-Goñi**, Zikun Chen, Sihai D Zhao, Winston Tan, Richard Weinshilboum, Krishna R Kalari, Liewei Wang  
*Cancer research* 83.8 (2023) pp. 1361–1380

## Conferences:

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### PRESENTED

Overcoming resistance in mCRPC: drug discovery using PDX models and high-throughput drug screening  
**Irene Marín-Goñi**, Huanyao Gao, Joachim L. Petit, Shreya Indulkar, Adam M. Kase, Cassandra N. Moore, Krishna R. Kalari, Michael T. Barret, Richard Weinshilboum, Wiston Tan, John A. Copland III, Liewei Wang  
*Poster presented in the American Association of Cancer Research (AACR Meeting)* (2025)

Targeting of CDK12/13 and androgen pathways in metastatic castration-resistant prostate cancer  
**Irene Marín-Goñi**, Shreya Indulkar, August John, Huanyao Gao, Mikel Hernaez, Liewei Wang  
*Poster presented in the American Society for Clinical Pharmacology and Therapeutics (ASCPT Meeting)* (2025)

Identification of transcriptional network disruptions associated to drug resistance in cancer with TraRe  
Charles Blatti, Jesús de la Fuente, Huanyao Gao, **Irene Marín-Goñi**, Zikun Chen, Sihai D Zhao, Winston Tan, Richard Weinshilboum, Krishna R Kalari, Liewei Wang, Mikel Hernaez  
*Poster presented in the European Conference of Computational Biology (ECCB)* (2022)

### ACCEPTED

Role of CDCP1 in Cardiac Fibrosis: spatially resolved gene expression profile in a Pressure Over-Load Mouse Model  
**Irene Marín-Goñi**, Li Wang, Min Wang, Rachad Ghazal, Will Sherman, Hridyanshu Vyas, Carolyn Roos, Runqing Huang, August John, Tamas Ordog, Mathew Wang, Liu Duan, Laura Lambert, Richard Weinshilboum, Naveen Pereira  
*Abstract accepted for poster presentation to American College of Cardiology (AAC) Meeting* (2025)

Discovery of a novel compound target CDCP1 to modulate human cardiac fibrosis  
Rachad Ghazal, Manikandan Selvaraj, Wang Min, Zhang Heyu, **Irene Marín-Goñi**, Mathew Coban, Caleb Weber, Naeyma Islam, Jessica Abbott, Duan Liu, Laura Lambert, Richard Weinshilboum, Thomas Caulfield, Naveen Pereira  
*Abstract accepted for poster presentation at the Basic Cardiovascular Sciences scientific sessions (BCVS) of American Heart Association (AHA)* (2025)

Interpretable Causal Representation Learning for Biological Data in the Pathway Space  
Jesús de la Fuente, Robert Lehmann, Carlos Ruiz-Arenas, **Irene Marín-Goñi**, Jan Voges, Xabier Martinez de Morentin, David Gomez-Cabrero, Idoia Ochoa, Jasper Tegnér, Vincenzo Lagani, Mikel Hernáez  
*Poster accepted at AIDrugX (NeurIPS)* (2024)

Encoding gene expression into gene set activity scores using neural networks  
Carlos Ruiz-Arenas, **Irene Marín-Goñi**, Idoia Ochoa, Mikel Hernaez, Luis A Pérez-Jurado  
*Poster accepted at Machine Learning for Computational Biology (MLCB)* (2023)

Bayesian Machine Learning Enables Identification of Transcriptional Network Disruptions Associated with Drug-Resistant Prostate Cancer  
Charles Blatti, Jesús de la Fuente, Huanyao Gao, **Irene Marín-Goñi**, Zikun Chen, Sihai D Zhao, Winston Tan, Richard Weinshilboum, Krishna R Kalari, Liewei Wang  
*Oral and Poster accepted at International Society for Computational Biology (ISCB), TransMed* (2021)