# Cristian Gonzalez-Colin

PhD Student, Bioinformatics and Systems Biology

 $Under graduate\ Intership$ 

Faculty of Medicine-INMEGEN

Updated October 3, 2023

CONTACT	<b>in</b> cris-gonzalezcolin <b>⊕</b> cristian2420.github.io	<b>⇔</b> cristian2420 <b>≥</b> cgonzalez@lji.org	$\bigcirc 0000$ -0002-3920- $\bigcirc cgonzalezcolin@u$	· .	JI ICSD
Research Interest	Understanding the effect of genetic variants linked to human diseases in immune-related cell types, through the development of computational tools.				
EDUCATION	University of California, San Diego			La Jolla, California	
	PhD in Bioinformatics and Systems Biology  Center for Genomic Sciences, UNAM  BS in Genomic Sciences  • Dissertation: Effects of disease risk-variants in gene expression			In Progress Morelos, Mexico 2016-2020	
	<ul> <li>at single cell level</li> <li>Thesis Comittee: Pandurangan Vijayanand, MD, PhD; Benjamin Schmiedel, PhD; Yvonne Rosenstein, PhD.</li> <li>Global Average: 9.3 out of 10</li> </ul>				
	Faculty of Sciences, UNAM BS in Biology			Mexico City, Mexico 2015-2016	
RESEARCH EXPERIENCE	PhD Student Vijayanand Lab, La Jolla Institute for Immunology Mentor: Pandurangan Vijayanand, MD, PhD			2022 – Present La Jolla, California	
	La Jolla Institute for Immunology  Research Technician			2019-2022	
	Vijayanand Lab, La Jolla Institute for Immunology Quantitative Trait Loci (QTLs) for gene expression			La Jolla, Ca	lifornia
	at bulk and single cell level, and histone marks in DICE database.  International Laboratory for Human Genome Research  Undergraduate Researcher			201	18-2019
	Regulatory Genomics and Bioinformatics Lab  Development of tools for the identification of conserved regulatory			Queretaro,	Mexico
	regions in Prokaryotes genomes for the RSAT suite tools.  Center for Genomic Sciences, UNAM			201	17-2019
	Undergraduate Researcher Computational Genomics Lab Development of machine learning tools for the improvement and			Morelos,	Mexico
	<ul> <li>automatization of analysis and biocuration on the REGULONDB.</li> <li>Automatic summarization of transcription factors (TFs) properties from text literature.</li> </ul>				
	<ul> <li>Supervised learning and text mining to retrieve regulatory interactions in bacterial literature.</li> <li>Text mining to retrieve transporter-substrate interactions.</li> </ul>				
	INMEGEN			201	18-2019

Analysis of preterm birth genomic markers in Mexican population.

Determination of cytokine concentration in preterm birth samples.

2015-2016

Mexico City, Mexico

#### Publications

- Pagadala, M., Sears, T. J., Wu, V. H., Pérez-Guijarro, E., Kim, H., Castro, A., Talwar, J. V., Gonzalez-Colin, C., Cao, S., Schmiedel, B. J., Goudarzi, S., Kirani, D., Au, J., Zhang, T., Landi, T., Salem, R. M., Morris, G. P., Harismendy, O., Patel, S. P., Alexandrov, L. B., Mesirov, J. P., Zanetti, M., Day, C.-P., Fan, C. C., Thompson, W. K., Merlino, G., Gutkind, J. S., Vijayanand, P., Carter, H., "Germline modifiers of the tumor immune microenvironment implicate drivers of cancer risk and immunotherapy response". eng. In: Nature communications 14.1 (May 2023), p. 2744. DOI: 10.1038/s41467-023-38271-5. PMID: 37173324.
- Schmiedel, B. J., Gonzalez-Colin, C., Fajardo, V., Rocha, J., Madrigal, A., Ramírez-Suástegui, C., Bhattacharyya, S., Simon, H., Greenbaum, J. A., Peters, B., Seumois, G., Ay, F., Chandra, V., Vijayanand, P., "Single-cell eQTL analysis of activated T cell subsets reveals activation and cell type-dependent effects of disease-risk variants". eng. In: Science immunology 7.68 (Feb. 2022), eabm2508. DOI: 10.1126/sciimmunol.abm2508. PMID: 35213211.
- Schmiedel, B. J., Rocha, J., **Gonzalez-Colin, C.**, Bhattacharyya, S., Madrigal, A., Ottensmeier, C. H., Ay, F., Chandra, V., Vijayanand, P., "COVID-19 genetic risk variants are associated with expression of multiple genes in diverse immune cell types". eng. In: *Nature communications* 12.1 (Nov. 2021), p. 6760. DOI: 10.1038/s41467-021-26888-3. PMID: 34799557.
- Chandra, V., Bhattacharyya, S., Schmiedel, B. J., Madrigal, A., **Gonzalez-Colin, C.**, Fotsing, S., Crinklaw, A., Seumois, G., Mohammadi, P., Kronenberg, M., Peters, B., Ay, F., Vijayanand, P., "Promoter-interacting expression quantitative trait loci are enriched for functional genetic variants". eng. In: *Nature genetics* 53.1 (Jan. 2021), pp. 110–119. DOI: 10.1038/s41588-020-00745-3. PMID: 33349701.
- Méndez-Cruz, C.-F., Blanchet, A., Godínez, A., Arroyo-Fernández, I., Gama-Castro, S., Martínez-Luna, S. B., **Gonzalez-Colin, C.**, Collado-Vides, J., "Knowledge extraction for assisted curation of summaries of bacterial transcription factor properties". eng. In: *Database: the journal of biological databases and curation* 2020 (Dec. 2020). DOI: 10.1093/database/baaa109. PMID: 33306798.

## Conference Presentation

Poster presentation at Keystone Symposia: Gene Regulation: From Emerging Technologies to New Models.

The cis-regulatory lanscape reveals cell type- and context-depedent effects of disease-risk variants affecting human immune cell types.

MENTORSHIP

### Elizabeth Marquez-Gomez

2021-2023

Summer 2022

Undergraduate Student, UNAM

La Jolla, California

Vijayanand Lab, La Jolla Institute for Immunology

Teaching

#### Teaching Assistant

Spring 2019

Center for Genomic Sciences

Morelos, Mexico

Bioinformatics Course.

Professors: Julio Collado-Vides and Heladia Salgado

OUTREACH

## Volunteer

Summer 2023

Camp Connect Science Class

CERTIFICATIONS

#### Introduction to Deep learning, UAEM

Fall 2015

SKILLS

Programming Languages: R, Python, Bash

Languages: English, Spanish