SEBASTIÁN AYALA-RUANO

I am a focused and motivated young researcher that has worked in **Bioinformatics** and Cheminformatics for four years at different laboratories. Phylogenetics and Structural Bioinformatics have been the main topics of this research experience. However, my current interests are devoted to Network Science, Complex Systems, and Machine Learning for drug discovery. Moreover, I am involved in various initiatives to empower Bioinformatics in Ecuador and Latin America.



EDUCATION

2020 2016

B.Eng., Biotechnology

Universidad San Francisco de Quito (USFQ)

Quito, Ecuador

- · Minor: Systems engineering.
- GPA: 3.78/4 (Magna Cum Laude) second best score of the College of Biological and Environmental Sciences 2020 class.
- Thesis: Computational study of two mutations in the Cereblon protein using molecular dynamics and quantum mechanics simulations (See details here).
- · Advisor: Miguel Angel Méndez Silva



RESEARCH EXPERIENCE

2021

Applied Signal Processing and Machine Learning Research Group - USFQ Quito, Ecuador Research Assistant

- We created a new method based on network science and similarity searching to discover new potential antiparasitic peptides (APPs). In brief, we explored the APPs' chemical space with three types of networks, predicted new potential leads using our best models, and found conserved motifs shared by our leads (See details here).
- Advisors: Yovani Marrero-Ponce, Noel Pérez Pérez

2020 2017

Computational and Theoretical Chemistry Group - USFQ

Undergraduate Researcher

Quito, Ecuador

- Identified binding specificity between repressor proteins and a transcriptional factor associated with the jasmonic acid pathway in Arabidopsis thaliana through molecular dynamics simulations and machine learning algorithms (See details here).
- I developed my undergraduate thesis in this laboratory.
- Advisor: Miguel Angel Méndez Silva

2020 2018

Bio-Chemoinformatics Group - Universidad de Las Américas

Research Intern

Quito, Ecuador

- Understood the impact of horizontal gene transfer in the genome of *Streptomyces* clavuligerus using phylogenetics, RNAseq data, and other bioinformatics tools (See
- Proposed molecular mimicry between Zika envelope protein and human neuronal proteins through structural similarity predictors, molecular dynamics, and proteinprotein interaction networks.
- · Advisor: Vinicio Armijos



CONTACT

- Sayalaruano.github.io
- ≤ sebasar1245@gmail.com











SKILLS

□ Technical

Languages:









DevOps:







Data Science:







Databases and Cloud:







A■ Languages

Spanish: Native English: Advanced Korean: Basic

> Made with the R package pagedown.

The source code is available at sayalaruano/cv.

Last updated on 2022-02-11.

View this CV online at sayalaruano.github.io/cv

Tumor Metabolism and Therapeutic Oncology Laboratory - Gwangju Institute of Science and Technology 2019

Research Intern

• Gwangju, South Korea

- · Performed a computational study of a missense mutation from Cereblon protein using molecular dynamics and quantum mechanics simulations, and other bioinformatics tools (See details here).
- Advisors: Miguel Angel Méndez Silva, Steve K. Cho

TEACHING EXPERIENCE

Linux and Bash/AWK scripting Boot Camp 2021

RSG Ecuador and iGEM Ecuador

♥ Virtual event

- Co-organizer and co-instructor. I designed and taught most of the course material.
- This course covered the basics of Linux, terminal usage, text and file processing command line tools, Bash/AWK scripting with applications in Bioinformatics, and Git/GitHub.

Undergraduate Teaching Assistant 2020

Learning Center - USFQ

Virtual events

 Provided online mentorship of Biotechnology, Mathematics, and Systems Engineering subjects to undergraduate students that needed help.

Undergraduate Teaching Assistant 2018

General Biology Laboratory - USFQ

Quito, Ecuador

- · Graded reports, tests, and other homework from the course.
- Provided feedback and guidance to undergraduate students in topics of the course.

PUBLICATIONS

Exploring the Chemical Space of Antiparasitic Peptides and Discovery of New Promising Leads through a Novel Approach based on Network Science and Similarity Searching

Preprint

2016

2021

2020

2020

· Ayala-Ruano S, Marrero-Ponce Y., Aguilera-Mendoza L., Pérez N., Agüero-Chapin G., Antunes A., Aguilar A. ChemRxiv (2021). doi: 10.33774/chemrxiv-2021-tgv69.

4th ISCB Latin American Student Council Symposium: a virtual and inclusive experience during COVID19 times

Editorial journal article

• Castillo-Vilcahuaman, C., Valdivia C., Osorio-Mogollón C., Silva-Andrade, C., Puche, R., Ayala-Ruano, S., Cuesta-Astroz, Y., Parra, G. F1000Research (2020). doi: 10.12688/f1000research.28330.1.

The molecular basis of JAZ-MYC coupling, a protein-protein interface essential for plant response to stressors Peer reviewed journal article

- Oña-Chuquimarca, S., Ayala-Ruano, S., Goossens, Pauwels, L., Goossens, A., Leon-Reyes, A., & Méndez, M. A. Frontiers in Plant Science (2020). doi: 10.3389/fpls.2020.01139.
- This article was chosen to feature in the Frontiers in Plant Science 2020 highlights e-book collection. doi: 10.3389/978-2-88966-723-9.

A putative antimicrobial peptide from Hymenoptera in the megaplasmid pSCL4 of Streptomyces clavuligerus 2019 ATCC 27064 reveals a singular case of horizontal gene transfer with potential applications Peer reviewed journal article • Ayala-Ruano, S., Santander-Gordón, D., Tejera, E., Perez-Castillo, Y., & Armijos-Jaramillo, V. Ecology and Evolution (2019). doi: 10.1002/ece3.4924. CONFERENCE PRESENTATIONS The molecular basis of JAZ-MYC coupling, a protein-protein interface essential for plant response to stressors 2021 ♥ Virtual event 6th Brazilian Student Council Symposium: Omics and Data Science • Oral presentation (slides and recording in English). Modeling of protein-protein interaction and search for key residues by machine learning of the JAZ-MYC3 2020 complex of Arabidopsis thaliana ♥ Virtual event 5th RSG-Argentine Symposium of Young Researchers in Bioinformatics • Oral presentation (slides and recording in Spanish). In silico detection of horizontal gene transfer in Streptomyces clavuligerus 2020 ♥ Virtual event International Society for Computational Biology Student Council Webinar series • Oral presentation (slides and recording in English). In silico detection of an antimicrobial peptide (AMP) transferred horizontally from arthropods to bacteria 2019 Plbagué, Colombia 2nd RSG-Colombia Symposium of young researchers in Bioinformatics • Oral presentation (slides in Spanish). Modeling of Protein-protein interaction and search for key residues by machine learning of a protein complex 2019 in the jasmonic acid route in Arabidopsis thaliana Urcuquí, Ecuador XLIII National Biology Conference · Poster presentation in Spanish. Structural changes due to a mutation in Cereblon might be a cause for intellectual disability 2019 Global Intern Program at Gwangju Institute of Science and Technology · Poster presentation in English. HONORS AND AWARDS 2021 Best oral presentation award Virtual event 6th Brazilian Student Council Symposium: Omics and Data Science Innovation challenge For more data on labor informality award 2021 Virtual event Datalat, PNUD Ecuador, UN Women Ecuador, and the International Labour Organization • This competition searched for a technological solution to collect labor informality data in Ecuador. There were 39 proposals from 80 interdisciplinary teams (See details about the challenge here). • The economic award of all stages of the competition was \$4.300

Chancellor's Honor List 2020 Ouito, Ecuador Universidad San Francisco de Quito 2016 • This award recognizes students who have a GPA of 3.7/4 or higher. Third HPC Summer School Colombia: Bio and Data Science scholarship 2020 Virtual event CyberColombia • The scholarship covered registration expenses for the event. 2nd RSG-Colombia Symposium of young researchers in Bioinformatics travel award 2019 Olombia Ibagué, Colombia **RSG Colombia** • This award covered the travel expenses to attend the event. **Global Intern Program** 2019 Gwangju Institute of Science and Technology • The GIP awarded students with accommodation and a monthly stipend to cover living expenses for eight weeks. During this time, we were involved in a research project and received valuable training and mentoring. LEADERSHIP AND SERVICE Current **Regional Student Group Ecuador** Quito, Ecuador International Society for Computational Biology Student Council 2020 · Co-founder and current president of the RSG Ecuador. This group aims to create a long-lasting community of students and researchers residing in Ecuador that work on Bioinformatics. **HerrCompBioinfo** 2021 ♥ Virtual event Regional Student Group Ecuador and Open Life Science (OLS) program • I led this project, an open-source educational resource of computational tools for Bioinformatics enthusiasts written in Spanish. • During the OLS program, I learned how to create and manage open science and open source projects. 17th Student Council Symposium 2021 Virtual event International Society for Computational Biology Student Council • I was the fellowship committee chair and contributed to other organization tasks. Saturdays.AI Quito 2021 project leader 2021 Virtual event Saturdays.Al Quito • I led my group project, which was an early plant disease detector based on convolutional neural networks, trained to recognize two types of maize infectious diseases. (See details here). 4th ISCB Latin American Student Council Symposium 2020 ♥ Virtual event International Society for Computational Biology Student Council

Note: I have developed other personal projects related to data science, machine learning, drug discovery, and other topics (See details here).

• I contributed to the program and fellowships committees.