

SEBASTIÁN AYALA-RUANO

I am a young researcher and freelance data scientist. I have worked in **Bioinformatics** and **Cheminformatics** for **five years** at different laboratories. My current research interests are devoted to **Network Science** and **Machine Learning** for drug discovery. I am part of several research, open-science, and software development communities (ISCBSC, The Carpentries, Streamlit Creators, and Open Life Science). Moreover, I am involved in various initiatives to empower Bioinformatics in Ecuador and Latin America.



EDUCATION

- 2024
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2022
- **MSc, Systems Biology**
Maastricht University (UM) 📍 Maastricht, the Netherlands
 - **Scholarship:** UM Holland-High Potential Scholarship for students from outside the EU/EEA.
 - **B.Eng., Biotechnology**
Universidad San Francisco de Quito (USFQ) 📍 Quito, Ecuador
 - **Minor:** Software engineering.
 - **GPA:** 3.78/4 (Magna Cum Laude) - second best score of the College of Biological and Environmental Sciences 2020 class.

WORK EXPERIENCE

- Oct. 2022
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Apr. 2022
- **Data scientist consultant**
Universidad de Las Américas 📍 Quito, Ecuador
 - Created a curated database of 50,000 herbarium records from tropical forest species of the Americas using web scraping and the [Global Biodiversity Information Facility](#) API.
 - Developed machine learning classifiers to predict the phenological stages of the forest species using the herbarium records. The models had values greater than 90% on all the performance metrics, and they can be used to design conservation strategies of the tropical forest species.
 - **Research Assistant**
Applied Signal Processing and Machine Learning Research Group - USFQ 📍 Quito, Ecuador
 - Created a method based on network science and similarity searching to explore the chemical space of antiparasitic peptides and discover new drugs (See details [here](#)).
 - **Advisors:** [Yovani Marrero-Ponce](#), [Noel Pérez Pérez](#)
 - **Research Intern**
[Computational and Theoretical Chemistry Group](#) - USFQ 📍 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](#)).
 - **Advisor:** [Miguel Angel Méndez Silva](#)
 - **Research Intern**
[Bio-Cheminformatics Group](#) - Universidad de Las Américas 📍 Quito, Ecuador
 - Understood the impact of horizontal gene transfer in the genome of *Streptomyces clavuligerus* using phylogenetics, and other bioinformatics tools (See details [here](#)).
 - Proposed molecular mimicry between Zika envelope protein and human neuronal proteins through molecular dynamics and protein-protein interaction networks.
 - **Advisors:** [Vinicio Armijos](#), [Yunierkis Perez](#)

CONTACT

🔗 sayalaruano.github.io
✉ sebasar1245@gmail.com



SKILLS

🖥 Technical

Programming Languages:



DevOps:



Data Science:



Databases and Cloud:



🗣 Languages

Spanish: Native

English: Advanced | C1 | TOEFL iBT 109

Korean: Basic

German: Basic

The source code is available at sayalaruano/cv.

Last updated on 2023-01-16.

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

2019

Research Intern

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

📍 Gwangju, South Korea

- Performed density functional theory and molecular dynamics simulations to understand the impact of a mutation in the ZN domain of the CRBN protein (See details [here](#)).
- I continued working on this research project in my undergraduate thesis.
- **Advisors:** Miguel Angel Méndez Silva, Steve K. Cho

**TEACHING EXPERIENCE**

2021

Co-organizer and co-instructor of a Bioinformatics boot camp

RSG Ecuador and iGEM Ecuador

📍 Virtual event

- 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.

2020

Undergraduate Teaching Assistant

Learning Center - USFQ

📍 Virtual events

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

2018

2016

Undergraduate Teaching Assistant

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****Peer reviewed journal articles**

- **Ayala-Ruano S.**, Marrero-Ponce Y., Aguilera-Mendoza L., Pérez N., Agüero-Chapin G., Antunes A., Aguilar A. (2022). *Exploring the Chemical Space of Antiparasitic Peptides and Discovery of New Promising Leads through a Novel Approach based on Network Science and Similarity Searching*. **ACS omega**. doi: doi.org/10.1021/acsomega.2c03398. Preprint: doi.org/10.26434/chemrxiv-2021-tgv69-v2.
- Oña-Chuquimarca, S., **Ayala-Ruano, S.**, Goossens, Pauwels, L., Goossens, A., Leon-Reyes, A., & Méndez, M. A (2020). *The molecular basis of JAZ-MYC coupling, a protein-protein interface essential for plant response to stressors*. **Frontiers in Plant Science**, 11, 1139. doi: [10.3389/fpls.2020.01139](https://doi.org/10.3389/fpls.2020.01139). This article was included in the Frontiers in Plant Science 2020 highlights e-book. doi: [10.3389/978-2-88966-723-9](https://doi.org/10.3389/978-2-88966-723-9).
- **Ayala-Ruano, S.**, Santander-Gordón, D., Tejera, E., Perez-Castillo, Y., & Armijos-Jaramillo, V. (2019). *A putative antimicrobial peptide from Hymenoptera in the megaplasmid pSCL4 of Streptomyces clavuligerus ATCC 27064 reveals a singular case of horizontal gene transfer with potential applications*. **Ecology and Evolution**, 9 (5), 2602-2614. doi: [10.1002/ece3.4924](https://doi.org/10.1002/ece3.4924).

Editorial journal articles

- Osorio-Mogollon C, Grentzinger V, Olguin-Orellana GJ, **Ayala-Ruano S.**, et al. (2023). *ISCB Student Council Symposium 2021, a virtual global venue: challenges and lessons learned*. **F1000Research**, 12(50). doi: [10.12688/f1000research.129945.1](https://doi.org/10.12688/f1000research.129945.1).
- **Ayala-Ruano S.**, Hernandez, F., Ortega, A., Infante, D., Carrascal, D., Sánchez-Luquez, K., & Puche-Quiñonez, R. (2022). *Highlights of the 1st Ecuadorian-Venezuelan Symposium of Young Researchers in Bioinformatics (ISEVIB)*. **F1000Research**, 11(1086), 1086. doi: [10.12688/f1000research.125381.1](https://doi.org/10.12688/f1000research.125381.1).
- Castillo-Vilcahuaman, C., Valdivia C., Osorio-Mogollón C., Silva-Andrade, C., Puche, R., **Ayala-Ruano, S.**, Cuesta-Astroz, Y., Parra, G (2020). *4th ISCB Latin American Student Council Symposium: a virtual and inclusive experience during COVID19 times*. **F1000Research**, 9. doi: [10.12688/f1000research.28330.1](https://doi.org/10.12688/f1000research.28330.1).

**SELECTED PRESENTATIONS**

- *Contributing Guidelines and Codes of Conduct for Open Projects*. (2022). Expert talk. **6th cohort of Open Life Science** (See details [here](#)).
- *Exploring the chemical space of antiparasitic peptides and discovery of new promising leads through a novel approach based on network science and similarity searching*. (2022). Oral presentation. **International Society for Computational Biology Student Council Webinar series** (See details [here](#)).

- HerrCompBioinfo: An open-source educational resource of computational tools for Bioinformatics enthusiasts written in Spanish. (2022). Oral presentation. **4th cohort of Open Life Science Graduation** (See details [here](#)).
- The molecular basis of JAZ-MYC coupling, a protein-protein interface essential for plant response to stressors. (2021). Oral presentation. **6th Brazilian Student Council Symposium: Omics and Data Science** (See details [here](#)).
- In silico detection of horizontal gene transfer in *Streptomyces clavuligerus*. (2020). Oral presentation. **International Society for Computational Biology Student Council Webinar series** (See details [here](#)).
- Structural changes due to a mutation in Cereblon might be a cause for intellectual disability. (2019). Poster presentation. **Global Intern Program - Gwangju Institute of Science and Technology** (See details [here](#)).



HONORS AND AWARDS

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|-------------------|--|
| 2022 | <ul style="list-style-type: none"> ● UM Holland-High Potential scholarship
Maastricht University 📍 Maastricht, the Netherlands

 <ul style="list-style-type: none"> • The UM Holland-High Potential Scholarship programme offers 24 full scholarships of € 30,000 (including tuition fee waiver and monthly stipend) each academic year for highly talented students from outside the European Union (EU) who have been admitted to a master's programme at UM. |
| 2021 | <ul style="list-style-type: none"> ● Best oral presentation award
6th Brazilian Student Council Symposium: Omics and Data Science 📍 Virtual event |
| 2021 | <ul style="list-style-type: none"> ● “For more data on labor informality” innovation challenge award
Datalat, PNUD Ecuador, UN Women Ecuador, and the International Labour Organization 📍 Virtual event

 <ul style="list-style-type: none"> • 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). |
| 2020

2016 | <ul style="list-style-type: none"> ● Chancellor’s Honor List and Magna Cum Laude
Universidad San Francisco de Quito 📍 Quito, Ecuador

 <ul style="list-style-type: none"> • These awards recognize students who have a GPA of 3.7/4 or higher. |
| 2020 | <ul style="list-style-type: none"> ● Third HPC Summer School Colombia: Bio and Data Science scholarship
CyberColombia 📍 Virtual event

 <ul style="list-style-type: none"> • The scholarship covered registration expenses for the event. |
| 2019 | <ul style="list-style-type: none"> ● 2nd RSG-Colombia Symposium travel award
RSG Colombia 📍 Ibagué, Colombia

 <ul style="list-style-type: none"> • This award covered the travel expenses to attend the event. |
| 2019 | <ul style="list-style-type: none"> ● Global Intern Program scholarship
Gwangju Institute of Science and Technology 📍 Gwangju, South Korea

 <ul style="list-style-type: none"> • 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

2022 | <ul style="list-style-type: none"> ● Streamlit creators program
Streamlit 📍 Virtual

 <ul style="list-style-type: none"> • I have created several web applications with the Streamlit Python package and became part of the community of creators. |
| Current

2021 | <ul style="list-style-type: none"> ● Open Life Science (OLS) program
Open Life Science 📍 Virtual

 <ul style="list-style-type: none"> • Leader of the HerrCompBioinfo project during the 4th cohort of the OLS program. We created an open-source educational resource of computational tools for Bioinformatics enthusiasts written in Spanish. Also, I learned how to create and manage open science and open source projects. • I mentored a group in the 5th cohort of the OLS program to create a computer vision-based tool to improve cancer diagnosis in Cameroon. See details about this project here. • I gave a talk about contributing guidelines and codes of conduct for open projects in the 6th cohort of OLS program. |

Current
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2020

● **Regional Student Group (RSG) Ecuador**

International Society for Computational Biology Student Council

📍 Virtual

- 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.
- Co-chair of the [1st Ecuadorian-Venezuelan Symposium of Young Bioinformatics Researchers](#).
- Fellowship committee chair of the [17th Student Council Symposium](#).
- Contributed to the program and fellowships committees of the [4th ISCB Latin American Student Council Symposium](#).

2021

● **Saturdays.AI Quito 2021**

[Saturdays.AI Quito](#)

📍 Virtual event

- 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](#)).

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