

BIOGRAPHICAL SKETCH

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NAME: Gonzalez, Fabio A.

eRA COMMONS USER NAME (credential, e.g., agency login): FABIOGONZALEZ

POSITION TITLE: Graduate Student Research Assistant

EDUCATION/TRAINING *(Begin with baccalaureate or other initial professional education, such as nursing, include postdoctoral training and residency training if applicable. Add/delete rows as necessary.)*

| INSTITUTION AND LOCATION | DEGREE (if applicable) | Start Date MM/YYYY | Completion Date MM/YYYY | FIELD OF STUDY |
|-----------------------------------|---------------------------|-----------------------|-------------------------------|-----------------------------|
| Universidad Icesi, Cali, Colombia | B.S. | 07/2013 | 08/2019 | Chemistry |
| Universidad Icesi, Cali, Colombia | B.S. | 07/2013 | 08/2019 | Pharmaceutical Chemistry |
| University of Delaware | PHD | 08/2019 | 08/2026 (expected) | Biochemistry |

A. Personal Statement

My research studies are directed towards understanding the function of membrane proteins and protein complexes, by studying the molecular determinants of protein-to-protein interactions at the atomistic resolution level. During my undergraduate research, I studied DNA-to-protein interactions by molecular dynamics and protein docking simulations. Following this, I spent two years of my graduate studies conducting large-scale simulations of membrane lipids and analyzing the dynamic properties of lipids in biological membranes. The early exposure to computational chemistry familiarized me with experimental tools for integrative modeling, such as NMR restraints, cross-linking restraints, electron microscopy density maps, etc. Complementary to my previous experience, I decided to continue my PhD research in the experimental field where I am interested in understanding the cellular role of selenoproteins, proteins that contain the genetically encoded amino acid selenocysteine. By elucidating the structure of an enzyme acting in cellular stress, selenoprotein S, and its mechanism of binding to protein partners we work to decipher the function of this protein in different cellular mechanisms.

So far, I have gained experience in both experimental structural biology and biochemistry techniques and theoretical techniques. My list of technical skills includes working with Linux operating systems, programming languages (Python, Tk-Tcl, R), and simulation software (NAMD, Amber, Gromacs). In addition, my experimental skills encompass the ability to purify membrane proteins, characterization of biomolecules by size exclusion chromatography and intact mass spectrometry, *in vitro* pull-downs, *in vitro* cross-linking, western blotting, and preparing biological samples for structural studies such as X-ray crystallography, imaging by negative staining EM and structural characterization by Cryo-EM.

My early experience in science has motivated me to pursue a postdoctoral training in an area related to cryo-EM and structural biology. The skills earned in my current Ph.D. career fit both the experimental and computational techniques required to gain experience in novel cryo-EM techniques and methods to characterize proteins and larger protein complexes. A long-term vision for me is to take my knowledge and experience to a research center to train and teach users to use structural techniques to characterize proteins related to their research projects. To accomplish this goal, besides my technical skills I have been working on developing leadership skills, and during my Ph.D. career, I have mentored two students (one undergraduate and one graduate) in our research

group and supervised two rotation students. I plan to expand this mentorship through my Ph.D. career and postdoctoral experience. I believe in the importance of education and I want to make it a personal goal of mine to transfer my knowledge and experience to other people as part of my job.

Born and raised in Colombia, coming to the United States to pursue my Ph.D. career was my first experience as an underrepresented group member. Being here on a solo status has granted me a different perspective based on the opportunities and resources provided to me due to my early efforts as an undergraduate. My research experience in Colombia was very limited, due to the limited number of resources, and I opted to do computational chemistry despite having an interest in other experimental fields. Given the opportunity to complete my graduate studies in the USA, and the chance of being able to receive high-level education and cutting-edge research resources has created a strong sense of commitment to finish my studies and pursue my career goals. This notion has also created a responsibility in me to give other underrepresented individuals help of any kind, such as opportunities, advice, and training. By providing a platform to give underrepresented students opportunities in research and education, we are creating a stronger scientific community with a sense of solidarity for those with a lack of representation. My ultimate desire is to break down barriers and stereotypes for underrepresented groups and serve as a role model to encourage others to pursue their career goals.

B. Positions, Scientific Appointments, and Honors

Positions and scientific appointments

2024-present Member, American Heart Association

2022-2023 Member, American Society for Biochemistry and Molecular Biology

2019-2020 Member, Biophysical Society

2018-2019 Member, American Chemical Society

2019-present Graduate Research Assistant, Department of Chemistry and Biochemistry, University of Delaware

Honors

2024 Brennie Hackley Jr. Award for Excellence in Research, Department of Chemistry and Biochemistry, University of Delaware

2024 Research Supplement to Promote Diversity in Science, American Heart Association

2020 Computer in Science & Engineering 2020 Best Paper Award

2019 Cum Laude (B.S in Chemistry with minor in Biochemistry), Universidad Icesi

2019 Cum Laude (B.S in Pharmaceutical Chemistry), Universidad Icesi

2013-2019 Dean's List, Universidad Icesi

C. Contributions to Science

- **Understanding the dynamical properties of viral envelopes through molecular dynamics simulations.**

My early research involved the mechanistic characterization of full-scale viral lipid bilayers. Using molecular modeling and employing massive parallel computers, I investigated the physical properties of the HIV-1 viral membrane at the coarse-grained resolution level. Using high-performance computing we derived the transverse diffusion rates of lipids in a native environment, an important physical property of heterogeneous lipid bilayers to maintain equilibrium.

1. Segura CP, Katyal N, **González-Arias F**, Bryer AJ, Perilla JR, Hadden-Perilla JA. Coronavirus through Delaware's Computational Microscope. Delaware Journal of Public Health. 2020 July;6(2):6-9. PMID: 34467099
2. **Gonzalez-Arias F**, Reddy T, Stone JE, Hadden-Perilla JA, Perilla JR. Scalable analysis of authentic viral envelopes on FRONTERA. Computing in science & engineering. 2020 August; 22(6):11-20. PMID: 33510584

- **Characterization of selenoprotein S cellular functions.**

Our research group is interested in understanding the different cellular roles of membrane-bound enzymes. My focus is selenoprotein S, which takes place in the resolution of ER stress and signaling pathways. The lack of structural information on this selenoprotein prevents understanding the primary function of selenoprotein S in different cellular processes such as Endoplasmic Reticulum Associated Degradation Pathway, inflammation, viral replication, gene regulation, etc. Thus, our ongoing efforts are

directed at characterizing the binding mechanism of selenoprotein S to molecular partners through biochemical techniques and structural techniques. So far, we have summarized in a review the latest insights into the structure, interactome studies, and cellular roles of selenoprotein S. I have also contributed to studies elucidating the role of selenoprotein S in SARS-CoV-2.

1. Ghelichkhani F, **Gonzalez FA**, Kapitonova MA, Schaefer-Ramadan S, Liu J, Cheng R, Rozovsky S. Selenoprotein S: A versatile disordered protein. Archives of Biochemistry and Biophysics. 2022 November 30;731:109427. PMID: 36241082
2. Ghelichkhani F, **Gonzalez FA**, Kapitonova MA, Rozovsky S. Selenoprotein S interacts with the replication and transcription complex of SARS-CoV-2 by binding nsp7. Journal of Molecular Biology. 2023 April 435(8), 168008. PMID: 36773692
3. Serbynovskiy V, Wang J, Chua EY, Ishemgulova A, Alink LM, Budell WC, Johnston JD, Dubbeldam C, **Gonzalez FA**, Rozovsky S, Eng ET. CryoCycle your grids: Plunge vitrifying and reusing clipped grids to advance cryoEM democratization. bioRxiv. 2024 Jan 24:2024-01

Complete List of Published Work in My Bibliography:

<https://www.ncbi.nlm.nih.gov/myncbi/fabio.gonzalez.1/bibliography/public/>

D. Scholastic Performance

| YEAR | COURSE TITLE | GRADE |
|--------------------------|--|-------|
| Universidad Icesi | | |
| 2013 | Algebra | 4.5 |
| 2013 | Logic and Argumentation | 4.4 |
| 2013 | Written communication I | 3.6 |
| 2013 | General Biology and Lab | 4.3 |
| 2013 | General Chemistry and Lab | 4.5 |
| 2014 | Industrial Safety and Hygiene | 3.8 |
| 2014 | Calculus I | 4.1 |
| 2014 | Written communication II | 3.8 |
| 2014 | Cellular Biology and Lab | 4.5 |
| 2014 | Organic Chemistry I and Lab | 4.6 |
| 2014 | Introduction to Pharmaceutical Sciences | 4.3 |
| 2014 | Organizations | 4.0 |
| 2014 | Calculus II | 4.8 |
| 2014 | Physics and Lab | 4.3 |
| 2014 | Biochemistry and Lab | 4.3 |
| 2014 | Organic Chemistry II and Lab | 4.5 |
| 2015 | Biostatistics | 4.6 |
| 2015 | Physical Chemistry I and Lab | 4.3 |
| 2015 | Analytical Chemistry and Lab | 4.4 |
| 2015 | Experimental Design | 4.3 |
| 2015 | Differential Equations | 4.0 |
| 2015 | Microbiology and Lab | 3.9 |
| 2015 | Physical Chemistry II and Lab | 4.7 |
| 2015 | Instrumental Analysis and Lab | 4.4 |
| 2016 | Principles of Economy | 4.7 |
| 2016 | Principles of Constitutional Law | 4.9 |
| 2016 | Organic Chemistry III and Lab | 4.1 |
| 2016 | Pharmacognosy and Phytochemistry and Lab | 4.5 |
| 2016 | Pharmacotechnics I and Lab | 4.4 |

| YEAR | COURSE TITLE | GRADE |
|-------------------------------|--|----------|
| 2016 | Molecular Biology | 4.2 |
| 2016 | Biotechnology and Lab | 4.2 |
| 2016 | Anatomy and Human Physiology and Lab | 4.1 |
| 2016 | Principles of Bioinorganic | 4.6 |
| 2016 | Enzymology and Lab | 4.5 |
| 2017 | Inorganic Chemistry and Lab | 5.0 |
| 2017 | Computational Biochemistry | 4.1 |
| 2017 | Pharmacotechnics II and Lab | 4.1 |
| 2017 | Public Health and Management | 4.1 |
| 2017 | Pharmacology | 4.0 |
| 2017 | Nutrition, Bromatology and Lab | 4.5 |
| 2017 | Physical Chemical and Microbiological Control | 4.5 |
| 2017 | Pharmacokinetics and Lab | 4.5 |
| 2017 | Toxicology and Lab | 4.2 |
| 2017 | Industrial Pharmacy and Lab | 4.3 |
| 2017 | Introduction to Drug Discovery | 4.5 |
| 2018 | Industrial Microbiology and Lab | 4.5 |
| 2018 | Bioethics | 4.0 |
| 2018 | Advanced Biochemistry and Lab | 4.3 |
| 2018 | Statistical Mechanics | 4.5 |
| 2018 | Clinical Pharmacy | 4.3 |
| 2019 | Senior Project | Approved |
| University of Delaware | | |
| 2019 | Quantum Chemistry | A- |
| 2019 | Solid State NMR Spectroscopy | A |
| 2020 | Chemical Dynamics | A- |
| 2020 | High-Performance Computing for Scientific Applications | A |
| 2020 | Biophysical Chemistry | A |
| 2020 | Chemical Thermodynamics | A |

Universidad Icesi grades classes from 0 to 5. Passing is 3.0 or better.