

# LAUREN SABO

(678) 780 1401 | lsabo8@gatech.edu | linkedin.com/in/laurengsabo | github.com/laurengsabo

Hardworking, Computer Science student aiming to contribute skills in big-data, multiprocessing, software development, and application-based problem solving to the software industry as an intern.

## Education

**Georgia Institute of Technology**, Atlanta, GA

May 2025 (expected)

**Major:** Computer Science, Concentrations: Intelligence & People

- *Minor:* Health & Medical Sciences
- Dean's List at Georgia Tech

## Research

**McGrath & Streelman Lab**, *Bioinformatics Researcher*

May 2022 — Present

A population genomics lab with a focus on Lake Malawi Cichlids

- Programmed a multiprocessing script within Python for the Genome Analysis Tool Kit (GATK) to process big data, such as multiple FASTA and VCF files at once.
- Spearheaded the processing and analysis of large-scale genomic datasets for Lake Malawi cichlids, utilizing advanced bioinformatics tools and programming languages such as Python.
- Collaborated with graduate students to share resources, data, and methodologies, fostering a cooperative research environment, and contributing to the broader scientific community's understanding of cichlid evolution.
- Writing an undergraduate thesis on the genomic assembly and annotation of Lake Malawi cichlids, contributing to the generation of high-quality reference genomes to enhance understanding of population structure and evolutionary dynamics.

## Projects

**"Expanding the Lake Malawi cichlid genome using high quality long-read sequencing", Head Researcher**

Jan. 2024 — Present

- Executing rigorous bioinformatics analyses to assemble and annotate complex genomic sequences, enhancing the genomic resources available for studying cichlid speciation and diversification.
- Enhancing the existing *M. zebra* reference genome (UMD2a) and developing the reference genomes of *M. conophoros*, *C. virginalis*, *A. chitande*, and *L. fuelleborni*.

**Chattahoochee River Conditions App – iOS Development**, *Full Stack Software Engineer*

Jan. 2024 — Present

- Developing in React Native and JavaScript, the app is intended to provide real-time river conditions for the GT Rowing Club.
- Is connected to the USGS Watershed Services database for updates and tells the user when the conditions are unsafe/ safe.

**"An Alternative Microinjection Manipulator and Pipette Pulling Technique as a Tool for the Delivery of CRISPR-Cas9", Head Researcher**

Sept. 2020 — Apr. 2021

- Designed a cost-effective approach to pulling Microinjection needles and 3D-printed a micromanipulator.
- Apparatus and needles can be used for in-vivo injections into the gonad of the nematode *C. elegans*.

## Work Experience & Leadership

**GT Rowing Club**, *Rower, Coxswain, & Board Member*

Feb. 2022 — Present

*Head Coxswain (2024):* Leader of coxswain education and manager of coxswain equipment.

*Merchandise Officer (2023):* Developed a team brand and coordinated merchandise inventory with distributors.

- Medaled at the Head of the Hooch Regatta (Chattanooga, TN), Southern Intercollegiate Rowing (SIRA) (Knoxville, TN), Head of the South Regatta (Augusta, GA), Secret City Head Race (Knoxville, TN).
- Rowed in A-Finals at the Jefferson Dad Vail Regatta (Philadelphia, PA), twice.
- Requalified for the Head of the Charles 2022 (Boston, MA) as a coxswain in the Women's Collegiate 4+ event, placing 13th/ 35.
- Commits 25+ hours weekly to conducting and overseeing rowers, practice, and team building tactical assistance during races.

**Gold & Bold Coffee Roasters**, *Barista*

Nov. 2022 — Present

- Skilled in customer service and food prep in a high-speed environment.

## Skills & Interests

**Computing Languages & Software:**

Java, Python, JavaScript, React Native, C, R, Assembly Code, HTML, CSS, Docker, Swift

**Bioinformatics:**

*Dry-Lab:* Genome Sequencing Software (HiFi.asm, LJA, Flye), BioNano UI, Server Organization Strategies, Binary Alignment Map (BAM) Files, Variant Call Format (VCF) Files, Genome Analysis Tool Kit (GATK)

*Biotechnology:* DNA Isolation, Qubit Quantification, Nanodrop Quantification, Gel Electrophoresis, SDS-PAGE Gel Electrophoresis, Restriction Digest/PCR, ELISA, Care of BSL1-Level Bacteria, Stain Protocols, Care of *C. elegans*

**Concepts:** Object-Oriented Programming (OOP), Genomic Assembly, Computational Biology, Multiprocessing, Biotechnology, Front-End Software Development, Computer Architecture, Data Structures, Linear Algebra, Differential Equations

**Affiliations:** GT Rowing Club, Women @ the College of Computing, Little Einsteins Outreach Club

**Hobbies:** Hiking, Traveling, Drawing, Reading, Painting, and Architecture