

# Jacob C. Fuller

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## Summary

Python programmer working in UGA Dept. of Genetics lab performing computational analyses of next-gen sequences.

## Education

*University of Georgia*

*August 2011 - May 2015*

- B.S. Genetics, Certificate of Computing. GPA: 3.3

## Skills

- Java, OOP 1 year
- Linux 3 years
- Shell 1 year
- HTML, CSS, JavaScript < 1 year
- Python 1 year
- Git 1 year
- Cluster computing 1 year
- C & C++ < 1 year

## Experience

### **Research Technician - UGA**

*White lab, Dept. of Genetics*

*June 2016 - present*

- Conducting independent computational analyses of next-gen sequence data, using cluster computing and Python scripts (see github)
  - Identify relevant and high quality whole-genome sequences on SRA database
  - Use UGA GACRC high-performance cluster to download and run a series of genomics software in order to process large amounts of sequence data. Run parallel cluster jobs for efficiency
  - Write Python scripts to analyze patterns in sequence depth to elucidate Y chromosome structural variation
  - Wrote algorithm that successfully identifies the boundaries of pseudoautosomal regions in Stickleback fish
  - Write Python scripts to analyze SNP (single-nucleotide polymorphism) density and diversity
  - Ultimate goal of this project is to understand Stickleback Y chromosome diversity at the population level
  - Weekly meetings with PI to discuss progress and scientific literature
- Molecular genetics experiments
  - Use bacterial artificial chromosomes for DNA transformation, cloning, extraction, and purification for genome assembly
    - troubleshooted protocol to increase DNA yield
  - PCR & PCR cloning using *E. coli*

*Steet lab, Complex Carbohydrate Research Center*

*June 2015 - June 2016*

- Bred and maintained transgenic strains of zebrafish
  - Maintained transgenic strain purity by genotyping via fluorescent microscope. Each transgenic line had a fluorescent protein marker
- Performed zebrafish husbandry and facility maintenance. I maintained strict organization and scheduling in order to keep the fish healthy and the Aquaneering fish systems running smoothly

***Undergraduate Researcher***

*January - May 2015*

*Kissinger Lab, Center for Tropical and Emerging Global Diseases*

- Performed computational analyses of NUMTs (nuclear mitochondrial transfer sequences) in order to identify strain specific sequences in *Toxoplasma gondii*
- Wrote Java program to parse and filter .txt files of NUMTs and converted this data to a more human-readable excel sheet format
- Participated in weekly lab meetings to learn from the graduate students' projects and presentations

## **Volunteering**

***MEDLIFE***

*August 2011 - May 2014*

*Mobile Clinic trip to Lima, Peru*

*December 2011*

- Spent a week setting up medical tents with a group of volunteers and local Peruvian doctors to provide healthcare to people in the pueblo jóvenes surrounding Lima, serving upwards of 250 individuals a day
- Worked alongside locals in community development projects. Specifically, we constructed a large staircase in the steep, arid mountainsides where the locals lived

*Executive Board Member: Spring Benefit Concert Chair*

*May 2013 - May 2014*

- Organized and ran the annual battle of the bands benefit concert. Raised over \$1000 and beat the previous record
- Assisted with general MEDLIFE executive board tasks

***Grady Memorial Hospital Observership***

*June 2012 - August 2012*

- Shadowed attending physician and residents during the ER night shifts 11PM -4AM
- Saw a wide variety of emergency medical issue and assisted physicians as needed