

# Alexander Larsen

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## TECHNICAL SKILLS

### Multi-purpose Language

- *Python*
- *R*
- *C++*
- *JavaScript*
- *Bash scripting*

### Sublanguages

- *SQL*
- *MongoDB*
- *CSS / HTML*

### System / Support

- *Git*
- *AWS CLI*
- *REST API*
- *Docker*
- *GIMP img editor*

## SUMMARY

Statistical analyst of Next Generation Sequencing (NGS) data as it relates to clinical applications. Excels in both fast-paced environments with urgent clinical results and methodical high dimensional data analysis. Automates multi-language bioinformatic software and tool development with an emphasis on python. Works with business partners and other researchers coordinating projects and public speaking. A self-directed learner, highly collaborative, and excellent at working autonomously.

## ACHIEVEMENTS

- Designed automated COVID-19 report end to end software.
- Developed metagenomic analysis library that automatically pulls, graphs, and models metadata.
- Designed automated multi-process Human Microbiome Project Whole Genome Sequencing (WGS) metagenomic pipeline.
- Created rapid primer design tools for over 10,000 different microscopic species.
- Presented on bioinformatics clinical value to congress representative and at other local events.
- Published scientific article with first authorship.

## EXPERIENCE

### Clinical Bioinformatician II

*Aperiomics Inc.*

**June 2018 – Current**

*Sterling, VA*

- Pipeline software development and module creation. Python, SQL, command-line/bash, C++, HTML, CSS, Javascript and R • Quality control and standards development.
- NGS protocols optimization and data analysis. Statistical modeling, Bayesian, Regression, PCoA, diversity analysis.
- Public speaking, clinician consults, and patient customer service.
- Publication writing, SOP writing, and publication grade image editing.

### Undergraduate Researcher

*University of Wisconsin Madison*

**Apr 2013 – May 2016**

*Madison, WI*

- Characterizing circadian rhythm genes and their temporal effect on long term memory formation in drosophila. Particular effort towards the quality control of data and samples.
- Characterizing zebrafish embryonic development, crayfish nerve reading frequencies, and rat hippocampal electrical activity.

## EDUCATION / CERTIFICATIONS

### Bioinformatics Micromasters Certificate

*UMGC - edX*

**Mar 2019 – Jan 2020**

*Adelphi, MD*

- Proteins and DNA: Alignment, Statistical Analysis, and Structure. BIF001x - BIF003x

### Bioinformatics

*Audited – UW Madison / MIT Open Courseware*

**June 2017 – Apr 2018**

*De Pere, WI*

- Graduate studies following UW Madison outlines for a master's in bioinformatics.
- Statistics, bioinformatic algorithms I/II, general algorithms / data structures, artificial intelligence.

### B.S. Neurobiology

*University of Wisconsin Madison*

**Sept 2012 – May 2016**

*Madison, WI*

- Graduated from a selective honors biology program which facilitated group directed research, scientific writing, and analysis. – UW Madison Biocore program.