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## **EDUCATION**

#### MIDDLEBURY COLLEGE

B.A COMPUTER SCIENCE Expected Feb 2020 | Middlebury, VT Cum. GPA: 3.41/4.0

#### **CUNY HUNTER COLLEGE**

Sep 2014 - May 2015 | New York, NY Cum. GPA: 4.0 / 4.0

## LINKS

Github:// AumitLeon LinkedIn:// AumitLeon Twitter:// @aumitleon Medium:// AumitLeon

## COURSEWORK

### **UNDERGRADUATE**

Mathematical Foundations of C.S Data Structures Computer Architecture Theory of Computation Machine Learning Quantum Computing Algorithms

# SKILLS

#### **PROGRAMMING**

Expert:

C++ • Python • C • Matlab JavaScript • Łate Proficient: Java • Ruby • Bash Prior Experience:

R • MySQL • Postgres

#### FRAMEWORKS/LIBRARIES:

Node.js • React.js • Express • D3.js • Django • ggplot2

- NumPy Pandas TensorFlow
- Scikit-Learn PyTorch

#### **GENERAL TECHNOLOGIES:**

HTML • CSS • Bootstrap • Linux/UNIX • SQL • Git/Github • AWS

#### **OTHER INTERESTS/SKILLS:**

Data Science • Statistical/Machine Learning • Bioinformatics • Open Source • Systems

## **EXPERIENCE**

# INDIGO AGRICULTURE | Consulting Software Engineer

Sep 2017 - Current | Charlestown, MA

- Working to enhance, maintain, and improve Indigo's bioinformatics server platform.
- Creating more efficient research pipelines through the Galaxy open source bioinformatics ecosystem.
- Providing support for Data Sciences projects, including both R+D and commercial projects.
- Developing customized solutions for the R&D pipeline, with a focus towards open source software and AWS infrastructure.

### INDIGO AGRICULTURE | SOFTWARE ENGINEERING INTERN

July 2017 - Aug 2017 | Charlestown, MA

- Developed solutions that made informatics analysis more efficient for non-technical scientists within Indigo's Research & Development department.
- Developed, prototyped, and deployed a bioinformatics server that accommodated genomic and proteomic analysis through the use of existing tools popular in the wider quantitative biological sciences community (such as BLAST, ClustalW, Bowtie).
- Pushed a Galaxy instance to production while maintaing an active pipeline for the integration of in-house tools.

# **SQUID INC. (FLEEK)** | SOFTWARE ENGINEERING INTERN

Sep 2015 - Dec 2015 | New York, NY

- Maintained internal developer dashboards via Node. JS and Python.
- Optimized for growth of our user base by creating internal tools and managing our Parse database.
- Oversaw the launch of our app to 1000+ more campuses during my internship.

# WIEGANG QIU BIOINFORMATICS LAB | LAB INTERN & ASSISTANT

Jan 2013 - May 2013 | New York, NY

- Assisted in genomic analysis of the Lyme disease pathogen using Perl, and internal PostgreSQL databases.
- Used genomic data to reconstruct evolutionary pathways and prepared results for publication.

# **PROJECTS**

#### OBSERVING THE EVOLUTION OF MUSIC WITH THE MSD

Nov 2017 - Dec 2017

Worked with Mariana Echeverria to analyze the Million Songs Dataset. We used deep Neural Nets and K-NN clustering to predict the year a song was released (75% accuracy), and K-Means clustering for more granular analysis of music features.

#### **EMAIL SPAM CLASSIFICATION**

Oct 2017 - Nov 2017

Developed a highly accurate Support Vector Machine model that could classify emails from the preprocessed Enron Spam dataset as spam or ham (non-spam). Achieved 99.6% accuracy, best in class.