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OBJECTIVE: To obtain a full-time position in front-end Software Engineering.

## **EDUCATION**

Carnegie Mellon University Pittsburgh, PA

M.S. Computer Science (Computational Biology)

AUG 2014

MAY 2016
(EXPECTE)

Honor Department Merit Fellowship (Tuition Scholarship) Selected

Sun Yat-sen University Guangzhou, China

SEP 2010 JUN 2014

**Programming** B.S. Biological Science

Python Honor BEST SOFTWARE tool project and GOLD AWARD in 2013 iGEM (International Genetic Java Engineering Machine) world competition held in MIT, Boston (SYSU-Software team member)

**WORK EXPERIENCE** 

SQL **Philips** Briarcliff Manor, NY

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MAY 2015

Research Intern of Clinical Informatics Solutions and Services(CISS)

AUG 2015

R Developed an algorithm to analyze clinical data to identify genetic mutations caused for antibiotic resistance in bacterial pathogens in Python and validated the correlation of results by statistical

PHP methods.

CSS SELECTED PROJECTS

JavaScript

**Bootstrap** 

Illustrator

**SKILLS** 

Perl

С

Matlab Generalized Data Mining Platform

Shell Implemented all back-end machine learning algorithms such as Random Forest, Neural Network,

**Tools** Hidden Markov Model.

Git Social Network (Deployed on AWS EC2)

Hive Developed a dynamic website with several interactive features involving following stream, push

MapReduce wall, notification, message system.

Hadoop Design front-end visualization and UI based on Bootstrap framework and all logos and icons.

MySQL WebSnap Chatting Web Application

node is

**Web Development**Designed the UX and implemented the front end features based on Backbone.js of an real-time online private chatting application with disappearing messaging feature for multi-users using the

Parse PaaS as the backend.

jQuery Django

Biological/Clinical data mining and processing

backbone.js Analyze edited DNA sequence in two different human cell lines and estimated gene-editing

UI Design efficiency.

Photoshop Performed noise filtering, feature detection, object segmentation and simulation of

microphotographs by MatLab.

Sketch Visualized RNA and RNA-Binding proteins interactions network using R.