### CHI ZHANG

Cellphone:8646243413 · Email:czhang6@g.clemson.edu · Linkedin: chi-zhang-544599b3 LeetCode: https://leetcode.com/CZhang1223 · github: https://github.com/bakerston

#### **EDUCATION**

#### Clemson University SC,USA

Computer Science Readiness Sequence | Jan 2021 - Present M.S. in Computational Biophysics | Aug 2015 - May 2021

## **Peking University** Beijing, CHINA

B.S. in Physics | Aug 2011 - May 2015

#### **TECHNICAL SKILLS**

#### C/C++, Python, Linux, Windows, Shell, SQL, MATLAB, R

TRAINGING AND CERTIFICATIONS

# Software Engineering Training

Insight Workshop Academy Jan 2021 - Present

#### Statistics with R

Coursera | Oct 2020

### **Deep Learning Specialization**

Coursera | Sep 2020

# IBM Data Science Specialization

Coursera | Apr 2020

#### **HONORS**

#### Best defender

Academic Competition of School of Physics, Peking University | May 2013

#### 1st Place

Academic Competition of School of Physics, Peking University | May 2013

#### 2nd Prize

The 29th National Physics Competition for College Students | Dec 2012

#### **EXPERIENCE**

**Teaching Assistant** | Clemon University, USA | Aug 2015 - Jan 2018 , Jan 2019 - present

• Worked closely with the lead teacher to identify issues students are having and develop appropriate solutions, helped designing lab experiment

**Research Assistant** | Clemon University, USA | Jan 2018 - Jan 2019

• Performed computational simulation to study multiple biological systems

**Research Assistant** Peking University, CHINA | Aug 2013 - Aug 2014

#### **PROJECTS**

## **Prediction Model of the Selling Price of Homes** | Jul 2020 - Oct 2020

- Developed a model to predict the selling price of a given home
- Applied multiple model selection methods including AIC and BIC
- Evaluate the performance of the final model with 80+ independent variables
- Related Technologies: R, Bayesian model, Exploratory Data Analysis, Data Visualization

# Observation of Self-seeding Effect of Islet Amyloid Polypeptide (hIAPP) | Sep 2018 - Dec 2019

- Build a quantitative model to simulate hIAPP interaction with fibril
- Analyzed actionable insights of hIAPP based on  $\sim$ 1TB of collected data
- Showed a full picture of fibril growth, which helps understanding the amyloid protein fibrillization process
- Related Technologies: Python, MATLAB, Linux, Shell

## **Inhibition of aggregation of Amyloid NACore by Fullerenol** | Aug 2016 - Nov 2018

- Applied computational simulations using discreted molecular dynamics algorithm
- Underscore the surface chemistry to form hydrogen bonds, which helps the design of novel theranostics against amyloid diseases
- Published on Nanoscale with 10+ citations
- Related Technologies: Statistical Analysis, Time Series Analysis, Monte Carlo Simultaion, Linux, Shell

#### **PUBLICATIONS**

Sun, Y., Kakinen, A., **Zhang, C.**, Yang, Y., Faridi, A., Davis, T.P., Cao, W., Ke, P.C. and Ding, F., 2019. Amphiphilic Surface Chemistry of Fullerenols is Necessary for Inhibiting the Amyloid Aggregation of Alpha-synuclein NACore. Nanoscale, 11(24), pp.11933-11945.