# Yonghao Jin

(+86)152-5692-0794

jyh1@mail.ustc.edu.cn

USTC, 96th Jinghai Road, Hefei, 230026

## Education

## UNIVERSITY OF SCIENCE AND TECHNOLOGY OF CHINA (USTC), HEFEI, CHINA

Bachelor of Science in Biophysics, Department of Physics (Expected)

Sept. 2013 - June 2017

Overall GPA: 3.49/4.3 or 85.23/100

## **Open Source Projects**

#### **MMACLONE**

GitHub Repository: https://github.com/jyh1/mmaclone

A cross-platform term rewriting system (TRS) with syntax similar to *Wolfram Mathematica* which has received **over one hundred stars** on *GitHub*. This interpreter is turing complete, with sophisticate pattern matching facilities and featured with symbolic computation, lambda calculus, etc. It consists of thousands of lines of Haskell codes, including sufficient amounts of unit tests to ensure robustness. This project won me the *Featured Contributor* in the *Wolfram Community*.

#### **SEQUENCE**

GitHub Repository: <a href="https://github.com/jyh1/sequence">https://github.com/jyh1/sequence</a>

A windows GUI program developed with the GTK library. This program is mainly calibrated to help sequence validation in molecular cloning experiments. It has a user-friendly interface to generate fasta format file from related sequencing files which could be used for down-stream alignment validation.

# Research Experience

#### STRUCTURE AND SEQUENCE MOTIF OF RBP BINDING SITES - 2016-PRESENT

Advisor: Prof. Kun Qu, School of Life Sciences, USTC

- Redesign and rewrite an iCLIP-seq (individual-nucleotide resolution Cross-Linking and ImmunoPrecipitation) data analysing pipeline with Python
- Fine-tune an iCLIP peak-calling algorithm, greatly reducing the evaluation time in P-Value estimation
- Analyse PARS (parallel analysis of RNA structure) data and reconstruct transcript secondary structures from PARS probing data with ViennaRNA Package

 Integrate PARS and iCLIP data to detect RBP sequence and structure binding motif to better predict RBP binding sites

#### **VARIATION OF WATER MOLECULES VIBRATION WITH TEMPERATURE – 2015**

Advisor: Dr. Wei Zhao, School of Physics, USTC

- Peak detection and analysis on Raman spectrum of water molecules
- Design algorithm to remove Rayleigh backgrounds of Raman spectrum
- Analyse the change of molecules vibration from the peak shift in Raman spectrum and interpret the observation with variation in hydrogen bond strength with temperature

## Fields of Interest

Big data analytics; Data Mining; Bioinformatics; Biomedical informatics; Computational Biology;

## **Skills**

**Adept in:** Python, C/C++, Mathematica(Graphing, Programming), Haskell, Pascal, Scheme, Git, Atom

Familiar with: Bash, Matlab, C#, Java, JavaScript, HTML

## Standardised Tests

TOEFL iBT: 29(Reading) + 29(Listening) + 22(Speaking) + 27(Writing) = 107

**GRE General:** 170(QR) + 155(VR) + 3.5(AW)

## **Awards and Honours**

- Bronze Merit Scholarship 2014
- Bronze Merit Scholarship 2013
- First Prize, the 17th National Olympiad in Informatics in Provinces (NOIP) 2011