# Yi Zhang

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#### **EDUCATION**

Ph.D. candidate, Bioengineering and Biomedical Engineering

Expected May 2019

University of Illinois at Urbana-Champaign (UIUC)

Urbana-Champaign, IL

- Track: Bioinformatics and computational biology
- Dissertation: Functional interpretation of cancer-associated genetic variants.

**Bachelor of Science**, Biosciences

June 2014

University of Science and Technology of China (USTC)

Hefei, China

#### **RESEARCH EXPERIENCE**

Research Assistant, University of Illinois at Urbana-Champaign

2015 - Now

Functional interpretation of breast cancer-associated genetic variants.

- Developed integrative algorithms for large-scale and multi-omics NGS data analyses of cancer patients.
- Designed haplotype imputation-based statistics on RNA-seq data to reveal variants' allele-specific effects.
- Integrated resources such as TCGA, GTEx, 1000 Genomes, dbSNP, ENCODE for variant interpretation.
- Discovered biological functions of breast cancer-associated non-coding variants as modulators of transcription factor binding activity through cis-regulation by enhancers.

Machine learning models for gene regulation and cancer subtype prediction.

- Built Random Forest models for 3D chromosome regulation prediction using epigenomic data.
- Implemented Convolutional Neural Networks for breast cancer subtype prediction using genotype data.

Interactive web resource for machine learning analysis of biomedical data, and NGS techniques.

- Built web tools for clustering, dimensional reduction, and data visualization (Docker, Javascript).
- Built an interactive knowledge base of ~70 NGS techniques and analysis (3000+ views in 3 months).

## **Undergraduate Summer Researcher**, University of California at Los Angeles

2013

Developed signal detection algorithms and interfaces for neuromuscular data analysis and visualization.

## **PUBLICATIONS**

- Y. Zhang\*, M. Manjunath\*, S. Zhang, D. Chasman, S. Roy, and J.S. Song. "Integrative genomic analysis predicts causative cis-regulatory mechanisms of the breast cancer-associated genetic variant rs4415084." <u>Cancer Research</u>, 78(7), 1579-1591, (2018). \*co-first authors.
- **Y. Zhang**, M. Manjunath, Y. Kim, J. Heintz, and J.S. Song. "SequencEnG: an Interactive Knowledge Base of Sequencing Techniques." *Bioinformatics*, bty794, (2018).
- M. Manjunath, Y. Zhang, Y. Kim, S.H. Yeo, O. Sobh, N. Russell, C. Followell, C. Bushell, U. Ravaioli, and J.S. Song. "<u>ClusterEnG</u>: an interactive educational web resource for clustering and visualizing high-dimensional data." *PeerJ Comput. Sci.*, 4:e155, (2018).

#### **CONFERENCE AND POSTERS**

- Y. Zhang, M. Manjunath, S. Zhang, D. Chasman, S. Roy, and J.S. Song. "Integrative genomic analysis discovers the causative regulatory mechanisms of a breast cancer-associated genetic variant." *AACR Annual Meeting, Chicago, IL* (2018)
- **Y. Zhang**, M. Manjunath, and J.S. Song. "Identifying causal mechanisms of germline risk variants in breast cancer." *Bioengineering Graduate Student Symposium, UIUC* (2017)
- **Y. Zhang**, P. Gad, V. R. Edgerton. "Burst detection in EMG of stepping spinal cord injured rats." *Cross-disciplinary Scholars in Science and Technology summer research*, No.69, Los Angeles, CA (2013)

## SKILLS

**Programming**: Python, Linux/Bash, R, MATLAB, C, HTML/CSS, JavaScript, TensorFlow **Bioinformatics**: Biomedical data analysis, NGS data analysis, cancer genomics, statistics, machine learning

# **RELATED COURSES**

Computational Cancer Biology	Machine Learning	Multivariate Analysis
Stochastic Processes	Statistical Learning	Mathematical Statistics

# **COURSE PROJECTS**

Statistical Learning: Prediction of Movie votes and campus shooting with SVM and gradient boosting models.

Machine Learning: Classification of NGS technique articles using natural language processing and SVM models.

Stochastic Processes: Segmentation of copy number using HMM based on genotype data in cancer cells.

Statistical Data Analysis in Physics: A cross-entropy method for graph clustering and Cheeger constant estimation.

### PROFESSIONAL EXPERIENCE

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Invited Speaker, Bioengineering Graduate Seminar, UIUC	2018
NHGRI Short Course on NGS: Technology & Statistical Methods, Birmingham, AL	2016 2015-2018
Big Data to Knowledge (BD2K) KnowEnG center at UIUC	
Student study group organized and participated:	
Deep Learning Study Group	2018
Statistical Learning Study Group	2017
TEACHING EXPERIENCE	
Invited Speaker, Bioinformatics Seminar, <i>UIUC</i>	2016, 2018
<ul> <li>Presented bioinformatics lectures and lead discussions among graduate students</li> </ul>	
Teaching Assistant, Biomedical Instrumentation Lab, Bioengineering, UIUC	2014-2015
<ul> <li>Lead lab and discussions among graduate students, graded lab reports</li> </ul>	
HONORS AND AWARDS	

Best Poster Award, Bioengineering Graduate Student Symposium, UIUC	2017
Outstanding Student Scholarship, School of Life Sciences, USTC	2013-2014
Outstanding Undergraduate Student Honor, USTC	2013
UCLA Cross-disciplinary Scholars in Sci. and Tech. (CSST) Summer Research Scholarship, USTC	2013
Aegon-Industrial Foundation Scholarship, USTC	2012
Panasonic Elite Scholarship, USTC	2011

## **UNIVERSITY SERVICE**

Volunteer, Genome Day, Carl R. Woese Institute for Genomic Biology, UIUC	2017
Vice-chairman of Student Union, School of Life Sciences, USTC	2012-2013
Organizer, School Photography Exhibition	2012