# Tongqiu (Iris) Jia

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# **EDUCATION**

# • University of Washington (UW)

MPH in Epidemiology

Sept 2020 - Present

#### • University of California, San Diego (UCSD)

B.S. in Biology with Specialization in Bioinformatics Minor in Computer Science

 $Sept \ 2014 - Jun \ 2018$ 

#### **PUBLICATIONS**

#### • Papers

- Tongqiu Jia, Brenton Munson, Hana Lango Allen, Trey Ideker, and Amit R. Majithia. "Thousands of missing variants in the UK BioBank are recoverable by genome realignment." Annals of human genetics (2020).
- Daniel E. Carlin, Samson Fong, Yue Qin, Tongqiu Jia, Justin K. Huang, Bokan Bao, Chao Zhang, and Trey Ideker. "A fast and flexible framework for network assisted genomic association." iScience (2019).
- Sophie Aimon, Takeo Katsuki, Tongqiu Jia, Logan Grosenick, Michael Broxton, Karl Deisseroth, Terrence J. Sejnowski, and Ralph J. Greenspan. "Fast near-wholebrain imaging in adult Drosophila during responses to stimuli and behavior." PLoS biology (2019).
- o Justin K. Huang, **Tongqiu Jia**, Daniel E. Carlin, and Trey Ideker. "pyNBS: a Python implementation for network-based stratification of tumor mutations." *Bioinformatics* (2018).

#### • Posters

 Sophie Aimon, Tongqiu Jia, Terrence. J. Sejnowski, Ralph J. Greenspan. Network models of whole brain activity in behaving adult Drosophila. Program No.152.10.2018 Neuroscience Meeting Planner. San Diego, CA: Society for Neuroscience, 2018. Online.

#### RESEARCH EXPERIENCE

#### • Graduate Research Assistant, University of Washington

Dec 2020 – Present

Mentors: Prof. Janet Baseman

- o Participated the evaluation of the public health impact of exposure notification app WA Notify
- Performed analysis on statewide surveys of knowledge, attitude and perceptions of COVID-19
- Analyzed data on adaption and usage of WA Notify

#### • Application Programmer, Ideker Laboratory, UCSD

Aug 2018 – Aug 2020

Mentors: Prof. Trey Ideker, Dr. Daniel Carlin

- Implemented network-based GWAS method and corresponding software and web interface
- Developed network-based GWAS methods for cross-species translation and shared disease mechanism discovery
- Analyzed genotype array and whole exome sequencing data for variants discovery
- Undergraduate Research Assistant, Ideker Laboratory, UCSD

  Mentors: Prof. Trey Ideker, Dr. Daniel Carlin. Dr. Justin Huang
  - Participated Python software development of Network Based Stratification algorithm (pyNBS)
  - Characterized features and parameters of different network propagation algorithms

# • Lab Technician, Salk Institute for Biological Studies

Sept 2017 – Jun 2018

Mentors: Prof. Terrence Sejnowski, Dr. Sophie Aimon

- Processed and analyzed large fruit fly whole brain image and behavioral datasets
- Automated and optimized fruit fly whole brain imaging data and behavioral data analysis pipeline

#### TEACHING EXPERIENCE

• Graduate Teaching Assistant

DATA 512: Human-Centered Data Science

Instructor: Brandon Martin-Anderson

• Teaching Assistant

Pathway and Network Analysis for Omics Data

Instructor: Professor Ali Shojaie

Human Centered Design and Engineering, UW Sept 2020 - Dec 2020

Summer Institute in Statistical Genetics, UW

July 2020

# Additional Experience

- System Administrative Assistant, Recreation Department, UCSD Oct 2015 Apr 2017
  - o Troubleshot computer software issues and set up operating systems and software on new computers
  - o Maintained website (recreation.ucsd.edu) and published web page updates under supervision
- Volunteer, Hamilton Glaucoma Center, Shiley Eye Institute, UCSD May 2015 Oct 2016
  - o Performed data collection and database entry in accordance with established protocols
  - Interacted with patients prior to their appointment in preparation for their visits

#### SKILLS

- Programming Intermediate: Python, Unix, Git
- Programming Basic: Java, R, C++, Matlab, JavaScript, React, SQL
- Bioinformatics Tools: BWA, GATK, PLINK,
- Machine Learning: Scikit-learn, Tensorflow, PyTorch
- Image Processing: Scikit-image, Nilearn, ImageJ
- Language: English, Mandarin