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Education

Massachusetts Institute of Technology (MIT)

Ph.D. in Theoretical Chemistry/Computational Biophysics

University of California, Los Angeles (UCLA)

Cross-disciplinary Scholars in Science and Technology Program

University of Science and Technology of China (USTC)

B.S. in Chemical Physics

Cambridge, Massachusetts
Sept. 2016-Present
Los Angeles, California
July-Sept. 2015

Hefei, China

Sept. 2012-June 2016

Research Interests

Computational modeling of biological systems from proteins to human genome, with potential applications to protein structure-function relationship and rational drug design using computational chemistry and machine learning

Research Experiences

• Predictive modeling of 3D organization of individual human chromosomes with Prof. Bin Zhang (MIT)

Feb. 2017 - Mar. 2018

- Developed and applied Maximum Entropy Algorithm to enable predictive modeling of 3D chromosomal structure
- Coded up core energy force field in Python and performed large-scale molecular dynamics simulations to generate structural ensemble of chromosomes and conducted statistical data analysis on simulated data
- o Data-driven modeling of whole-nucleus 3D human genome

with Dr. Alejandro Reyes (Broad) and Prof. Bin Zhang (MIT)

April 2019 - Feb. 2020

- Developed efficient data-driven modeling approach by extending the Maximum Entropy Algorithm to a whole-nucleus scale and identified important mechanisms dictating genome organization
- Coded up an open-source software in Python with three other languages and opened up various new research directions on mechanistic investigations of genome structure
- o Whole-genome modeling of genome reorganization in tumorigenesis

with Martin Aryee Lab (HMS/Broad) and Bradley Berstein Lab (MGH/Broad)

Mar. 2019 - June 2020

- Applied whole-nucleus genome organization modeling on real-patient data and revealed for the first time a large-scale reorganization of 3D genome organization in colorectal tumor cells in a quantitative manner
- Led computational modeling part in a massive collaboration among Harvard, MIT, Broad Institute and MGH
- Deep learning-based characterization of chromatin folding landscape

with Dr. Wenjun Xie and Prof. Bin Zhang (MIT)

Feb. 2019 - Oct. 2019

- Applied variational autoencoder together with *in silico* polymer model systems to analyze the fluctuation and heterogeneity of chromatin structures generated by single-cell imaging experiments

Papers

*Equal contribution.

Yifeng Qi, Bin Zhang. Predicting three-dimensional genome organization with chromatin states. *PLoS Computational Biology*, 15.6 (2019): e1007024.

Yifeng Qi, Alejandro Reyes, Sarah E. Johnstone, Martin J. Aryee, Bradley E. Bernstein, Bin Zhang. Data-driven polymer model for mechanistic exploration of diploid genome organization. *Biophysical Journal*, in press.

Sarah E. Johnstone*, Alejandro Reyes*, **Yifeng Qi**, et al. A topological atlas reveals layers of genome reorganization in colorectal cancer, *Cell*, 182.6 (2020): 1474-1489.

Wenjun Xie, **Yifeng Qi**, Bin Zhang. Characterizing chromatin folding coordinate and landscape with deep learning, *PLoS Computational Biology*, https://doi.org/10.1371/journal.pcbi.1008262.

Liangqi Xie, . . ., **Yifeng Qi**, . . ., Robert Tjian, Zhe Liu. 3D ATAC-PALM: Super-resolution Imaging of the Accessible Genome. *Nature Method*, 17.4 (2020): 430-436.

Zhaowei Chu, ..., Yifeng Qi, ..., Bin Zheng. STAG2 regulates interferon signaling in melanoma via enhancer loop reprogramming. *Submitted*.

Tao Chen, Tao Chen, Sheng Chen, Yuwei Zhang, **Yifeng Qi**, Yuzhou Zhao, Weilin Xu, Jie Zeng. Catalytic Kinetics of Different Types of Surface Atoms on Shaped Pd Nanocrystals. *Angewandte Chemie International Edition*, 55 (2016): 1839-1843.

Yifeng Qi, Bin Zhang. Exploring the free energy landscape of nucleolar coalescence in human cell nucleus. *In preparation*.

Xingcheng Lin*, **Yifeng Qi***, Andrew Latham, Bin Zhang. Learning mechanisms of genome folding with maximum entropy optimization. *In preparation*.

Liangqi Xie*, Peng Dong*, **Yifeng Qi***, et al. Super-resolution Imaging Reveals 3D Structure and Organizing Mechanism of Accessible Chromatin. *In preparation*.

Talks and Posters

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o NSF-funded Genome Architecture and Function Workshop

MIT, June 2020

MIT Biophysics Retreat

Cape Code MA, Oct. 2019

o Annual Greater Boston Area Statistical Mechanics Meeting

Brandeis University, Oct. 2019

Models, Inference & Algorithms (MIA) Seminar
 According to the Professional Medical Me

Broad Institute, Mar. 2019
Brandeis University Oct 2018

Annual Greater Boston Area Statistical Mechanics Meeting
Annual Greater Boston Area Statistical Mechanics Meeting

Brandeis University, Oct. 2018 MIT, Oct. 2017

Posters

o The Biophysics Society (BPS) Annual Meeting

San Diego CA, Feb. 2020

Molecular Biophysics in the Northeast Meeting

Northeastern University, Nov. 2019

o Gordon Research Conference-Stochastic Physics in Biology

Ventura CA, Jan. 2019

MIT Biophysics Retreat

Cape Code MA, Oct. 2018

o Keystone Symposia-Chromatin Architecture and Chromosome Organization

Whistler (Canada), Mar. 2018

Selected Awards

o Biophysics Society Graduate Student Award

BPS, 2020

Lester Wolfe Fellowship

MIT, 2017/2020

o Guo Moruo Scholarship (the highest honor for USTC graduates)

USTC, 2015

o Cross-disciplinary Scholar Science & Technology Scholarship

UCLA, 2015

National Scholarship (top 1% nationwide)

P.R.China, 2014

o National Olympiad Competition in Chemistry 1st Prize

P.R.China, 2011

Skills

Languages

Python, C, Matlab, Fortran, Shell, LATEX

Software

LAMMPS, VMD, Gaussian09, Github

Package Development
Courses Statistica

opment DRAGON (Github link: https://github.com/ZhangGroup-MITChemistry/DRAGON) Statistical Mechanics, Statistical Physics Biology, Physical Biology, Machine Learning, Algorithms for

Inference, Physics in Immunology, Computational Biology, Quantum Mechanics

Teaching Experiences

• Principle of Chemical Science (TA)

MIT, Sept. 2016-June 2017 USTC, Sept.-Dec. 2015

Quantum Physics (TA)

Activities

Scientific Journal reviewer for *Quantitative Biology*; Co-host for *Greater Boston Area Theoretical Chemistry Lecture Series* **Extra-Curricular** MIT Rowing Club; MIT Rock Climbing Club