# Ailun (Ellen) Wang

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#### Education

**Boston College** Chestnut Hill, MA

Ph.D. in Physical Chemistry Aug. 2015 - June 2021

Areas of Expertise: Computational Biophysics, Python, Data Analysis, High Performance Computing, Predictive Model, Data Visualization.

University of Science and Technology of China (USTC)

Hefei, China

B. Sc. in Materials Chemistry

Aug. 2011 - July 2015

## **Professional Experience**

**Northeastern University** 

Boston, MA

Postdoctoral Fellow at Center for Theoretical Biological Physics (CTBP)

July 2021 - present

## Research Experience

Highlights: High-performance scientific computing; Large-scale data analysis; Advanced data visualization; Statistical modeling.

## Prediction of antigen recognition on T-cell receptors using Machine Learning-based classification

Northeastern University, CTBP

- Developed a physical model in Python for predicting binding specificity between T-cell receptors (TCR) and antigens.
- O Utilized a Machine Learning based method to classify contact interfaces in strongly vs. weakly binding TCR and antigens.
- o Achieved high transferability and sensitivity with 94% accuracy on blind test cases.

### Constructing 3D architecture of chromatin at high resolution

Northeastern University, CTBP

- Designed a transferable model for predicting 3D architecture of chromatin to investigate the structural features of gene.
- o Implemented physical models for molecular dynamics (MD) simulations in Python and optimized model parameters to align with experimental measurements.
- o Predicted the 3D architectures of chromatin segments that closely agree with experimental measurements and the manuscript is currently under review.

#### Demystify the effect of ions in conformational dynamics of ribosom using computational methods Boston College

- Developed a force field to facilitate MD simulations of large biomolecules (sub-million atoms) with ions.
- o Implemented an iterative parameter optimization protocol in Python and C++ for force field development.
- o Conducted long-timescale MD simulations on HPC clusters and analyzed structural features using statistical methods in Python from terabytes of raw data.
- o Provided the first computational explanation of how diffuse Mg<sup>2+</sup> and K<sup>+</sup> ions regulate large-scale conformational rearrangements of ribosome.
- o Published 6 peer-reviewed journal articles and 1 book chapter and presented at 8 conferences.

## Theoretical investigation of ion transportation mechanism in porous materials

Boston College

- o Illustrated the ion transport mechanism in porous materials using density functional theory (DFT) calculations and ab initio MD simulations in multi-disciplinary collaboration with materials science experimentalists.
- Published 2 peer-review journal articles.

# **Selected Honors and Awards**

2020: Engelhard Pingree Research Fellowship

Boston College

2019: IHPCSS Fastest CPU Code International HPC Summer School (IHPCSS) 2019

2013: Outstanding Student Scholarship USTC

## Technical Skills

Programming: Python (7 years), Shell Script, SQL, Mathematica, Matlab, C++, LATEX

Softwares & Tools: Linux/Unix, Git, HPC, OpenMP, MPI, Jupyter notebook, Pandas, NumPy, SciPy, Matplotlib, Seaborn, Plotly