

Ye Ding

+86 185 2533 5631 | dingye@westlake.edu.cn | github.com/dingye18

EDUCATION

Westlake University

Doctor of Computational Biophysics, School of Life Sciences.

Hangzhou

Sep 2018 – Mar 2024

Dalian University of Technology

Bachelor of Applied Physics, School of Physics.

Dalian

Sep 2014 – July 2018

PROJECTS

DP/MM | *Deep Potential, CHARMM36m, OpenMM, ORCA*

March 2020

- Integrated Deep Potential with the classical force field CHARMM36m to develop the hybrid force field DP/MM, which enhanced the accuracy of atomic forces from MM to QM levels in an **adaptively changing** DP region.
- DP/MM achieves greater **accuracy** than MM and offers more **efficiency** and **flexibility** compared to QM/MM.
- Enhanced the geometry of zinc coordination structures in zinc-protein interactions using DP/MM, achieving significant improvements over traditional MM approaches.

logP Prediction with Drude Force Field | *Drude, logP, FEP, CHARMM*

Dec 2018 – Oct 2019

- Calculated the octanol-water partition coefficient for small molecules by using free energy perturbation (FEP) and the Drude polarizable force field.

Binding Free Energy for Ca^{2+} and Calmodulin EF-Hands | *Drude, CHARMM36m*

Nov 2019 – Oct 2021

- Calculated the absolute binding free energy of Ca^{2+} to Calmodulin EF-hands using the Drude and CHARMM36m force fields.

EXPERIENCE

Award

- National Scholarship, 2018-2019
- First Class Award of Asia Supercomputer Challenge 2017

DP Technology Research Intern

Aug 2022 - Oct 2022

- Engaged in the development of Induced Fit Docking tools and the deployment.

DP Technology Researcher in CADD

Apr 2024 - Now

- Specializing in advancing Free Energy Perturbation tools, with a focus on implementing Hamiltonian Replica Exchange Molecular Dynamics (H-REMD) and Grand Canonical Monte Carlo (GCMC) methods.

PUBLICATIONS

- Ye Ding**, Jing Huang et.al, 2020. Predicting partition coefficients of drug-like molecules in the SAMPL6 challenge with Drude polarizable force fields. 10.1007/s10822-020-00282-5
- Qiaozhu Tan, **Ye Ding**, Jing Huang, 2021. Binding Energy and Free Energy of Calcium Ion to Calmodulin EF-Hands with the Drude Polarizable Force Field. 10.1021/acsphyschemau.1c00039
- Ye Ding**, Kuang Yu, Jing Huang, 2023. Data science techniques in biomolecular force field development. 10.1016/j.sbi.2022.102502
- Ye Ding**, Jing Huang. DP/MM: A Hybrid Model for Zinc-Protein Interactions in Molecular Dynamics. 10.1021/acs.jpcllett.3c03158
- Ye Ding**, Jing Huang. Implementation and Validation of an OpenMM Plugin for the Deep Potential Representation of Potential Energy. 10.3390/ijms25031448
- Zilin Song, **Ye Ding**, Jing Huang. Constant Advance Replicas Method for Locating Minimum Energy Paths and Transition States. 10.1002/jcc.27178

TECHNICAL SKILLS

Languages: Python, C/C++, CUDA, BASH

Software: OpenMM, DeePMD-kit, CHARMM, ORCA, GROMACS, DMFF, TensorFlow, PyTorch

Developer Tools: Git, VS Code, SLURM, PBS