Sikao Guo

■ sikaoguo@gmail.com | 🏫 https://sikaoguo22.github.io/ | 🗣 Baltimore, MD | 📵 0000-0002-7680-8060 | 🖪 Sikao-Guo

Education _

Institute of Physics, Chinese Academy of Sciences

Beijing, China

Ph.D. in Condensed Matter Physics

Sep. 2014 - Dec. 2019

Research Advisor: Professor Ping Xie; Thesis: "Studying the movement mechanism of dimer kinesin through theoretical modeling and numerical simulation"

Nankai University Tianjin, China

B.S. in Physics

Sep. 2010 - Jun. 2014

Experience _

Johns Hopkins University

Baltimore, MD

Postdoctoral Research

Jan. 2020 - present

Research Advisor: Professor Margaret E. Johnson

Publications

- [1] **Sikao Guo**, Ipsita Saha, Saveez Saffarian, Margaret E Johnson (2023) "Structure of the HIV immature lattice allows for essential lattice remodeling within budded virions," *eLife*, 12:e84881. [Link]
- [2] Qi Xie, Sea On Lee, Nitya Vissamsetti, **Sikao Guo**, Margaret E. Johnson, Stephen D. Fried (2023) "Secretion-Catalyzed Assembly of Protein Biomaterials on a Bacterial Membrane Surface," *Angew. Chem. Int. Ed.*, e202305178. [Link]
- [3] Yian Qian, Daniel Evans, Bhavya Mishra, Yiben Fu, Zixiu Hugh Liu, **Sikao Guo**, Margaret E. Johnson (2023) "Temporal control by cofactors prevents kinetic trapping in retroviral Gag lattice assembly," *Biophysical Journal*. [Link]
- [4] Sikao Guo, Alexander J Sodt, Margaret E Johnson (2022) "Large self-assembled clathrin lattices spontaneously disassemble without sufficient adaptor proteins," *PLOS Computational Biology*, 18, e1009969. [Link]
- [5] Sikao Guo, Ping Xie (2020) "A common chemomechanical coupling model for orphan and conventional kinesin molecular motors," *Biophysical Chemistry*, 264, 106427. [Link]
- [6] Xiao-Xuan Shi, **Sikao Guo**, Peng-Ye Wang, Hong Chen, Ping Xie (2020) "All-atom molecular dynamics simulations reveal how kinesin transits from one-head-bound to two-heads-bound state," *Proteins: Structure, Function, and Bioinformatics*, 88 (4), 545-557. [Link]
- [7] **Sikao Guo**, Xiao-Xuan Shi, Peng-Ye Wang, Ping Xie (2019) "Run length distribution of dimerized kinesin-3 molecular motors: comparison with dimeric kinesin-1," *Scientific reports*, 9, 16973. [Link]
- [8] **Sikao Guo**, Xiao-Xuan Shi, Peng-Ye Wang, Ping Xie (2019) "Force dependence of unbinding rate of kinesin motor during its processive movement on microtubule," *Biophysical Chemistry*, 253, 106216. [Link]
- [9] **Sikao Guo**, Wei-Chi Wang, Peng-Ye Wang, Ping Xie (2019) "Force dependence of velocity and run length of kinesin-1, kinesin-2 and kinesin-5 family molecular motors," *Molecules*, 24 (2), 287. [Link]
- [10] Yi-Ben Fu, **Sikao Guo**, Peng-Ye Wang, Ping Xie (2019) "Dynamics of cooperative cargo transport by two elastically coupled kinesin motors," *The European Physical Journal E*, 42 (4), 1-13. [Link]
- [11] Ping Xie, **Sikao Guo**, Hong Chen (2019) "A generalized kinetic model for coupling between stepping and ATP hydrolysis of kinesin molecular motors," *International journal of molecular sciences*, 20 (19), 4911. [Link]
- [12] **Sikao Guo**, Xiao-Xuan Shi, Peng-Ye Wang, Ping Xie (2018) "Processivity of dimeric kinesin □ 1 molecular motors," *FEBS Open Bio*, 8 (8), 1332-1351. [Link]
- [13] Ping Xie, **Sikao Guo**, Hong Chen (2018) "ATP-concentration-and force-dependent chemomechanical coupling of kinesin molecular motors," *Journal of Chemical Information and Modeling*, 59 (1), 360-372. [Link]
- [14] Xiao-Xuan Shi, Yi-Ben Fu, **Sikao Guo**, Peng-Ye Wang, Hong Chen, Ping Xie (2018) "Investigating role of conformational changes of microtubule in regulating its binding affinity to kinesin by all □atom molecular dynamics simulation," *Proteins: Structure, Function, and Bioinformatics*, 86 (11), 1127-1139. [Link]
- [15] **Sikao Guo**, Peng-Ye Wang, Ping Xie (2017) "Dynamics of dimeric kinesins: Limping, effect of longitudinal force, effects of neck linker extension and mutation, and comparison between kinesin-1 and kinesin-2," *International journal of biological macromolecules*, 105, 1126-1137. [Link]
- [16] **Sikao Guo**, Peng-Ye Wang, Ping Xie (2017) "A model of processive movement of dimeric kinesin," *Journal of Theoretical Biology*, 414, 62-75. [Link]

Conference Presentations _____

Cell Bio 2022 Washington, DC, USA

Talk: Defects Within the HIV-1 Immature Lattice Support Dynamic Remodeling and Protease Dimerization,

Dec. 2022

[Link]

Biophysical Society 66th Annual Meeting

San Francisco, CA, USA

Poster: Large Self-assembled Clathrin Lattices Spontaneously Disassemble Without Sufficient Adaptor Proteins,

Feb. 2022

[Link]

Stochastic Physics in Biology (Gordon Research Conference)

Ventura, CA, USA

Poster: Self-assembled clathrin lattices spontaneously disassemble without sufficient links to the plasma membrane,

Oct. 2021

Biophysical Society 65th Annual Meeting

Virtual

Poster: Modeling Nucleation and Kinetics of Clathrin Assembly by Membrane Localization,

Feb. 2021

2020 Annual Meeting of the APS Mid-Atlantic Section

Virtual

Talk: Modeling Nucleation and Kinetics of Clathrin Assembly by Membrane Localization,

Dec. 2020

Academic Service _____

Peer Reviwer

 $The \ Journal\ of\ Physical\ Chemistry,\ PROTEINS:\ Structure,\ Function,\ and\ Bioinformatics,\ PLOS\ Computational\ Biology,$

International Journal of Molecular Sciences, Cells, Applied Sciences, Biophysica, Biochemistry and Biophysics Reports,

Jan. 2020 - Present

Entropy, Molecules, Biophysica, Medicina

Guest Editor

Biomedicines, Biology

Jan. 2023 - Present

References ____

• Prof. Margaret E. Johnson

Ph. D, Associate Professor, at School of Arts & Sciences, Johns Hopkins University, Baltimore, MD, USA

margaret.johnson@jhu.edu

· Prof. Ping Xie

Ph. D, Professor, at Institute of Physics, Chinese Academy of Sciences, Beijing, China

∠ pxie@iphy.ac.cn