ZHENGQING ZHOU

Department of Biomedical Engineering, Duke University, NC, USA +86 18201100514 | email: johnkingzhou@gmail.com

EDUCATION

Peking UniversityBeijing, ChinaBachelor of Science, PhysicsSep 2017 – Jul 2021

University of California, Los Angeles

CA, USA

Exchange student at the Physics and Astronomy Department

Sep 2019 - Dec 2019

PUBLICATIONS

Zhou, Z. et al. Dynamics of SARS-CoV-2 and Host Immunity in Infection and Vaccine Protection. Under review at Nat. Commun. DOI: 10.21203/rs.3.rs-590500/v1

RESEARCH EXPERIENCE

Plasmids by the Numbers

Duke University

Advisor: Prof. Lingchong You, Department of Biomedical Engineering

Sep 2021 – Feb 2022

- Constructed an online database for plasmid dynamics in microbial community. (<u>www.plasmidsbynumbers.com/</u>)
- Collected ~800 entries of qualitative (e.g. antibiotic resistance) and quantitative data (e.g. transfer rate) of over 140 plasmids.

Niche Partitioning Shapes Plasmid Dynamics

Duke University

Advisor: Prof. Lingchong You, Department of Biomedical Engineering

Sep 2021 - Feb 2022

- Niche partitioning decouples interspecific interactions and shapes the interaction matrix into a partitioned one.
- Nutrient niche partitioning retains the conjugation matrix dense, while spatial niches partition the matrix.
- Fundamental difference between the interaction matrix and conjugation matrix shape leads to increased conjugation frequency, suggesting diverse nutrient supply increases plasmid abundance.

Dynamics of SARS-CoV-2 Infection and Host Immunity and Vaccine Protection

Peking University

Advisor: Prof. Fangting Li, School of Physics, Center for Quantitative Biology

Feb 2020 - Present

- Improved upon previous viral infection models by including the full immune system dynamics of innate, cellular and humoral immune cells, cytokines and antibodies.
- Derived a collective metric (immune efficacy) of innate and adaptive immune cells to quantify the strength of dynamic immune response and analyzed COVID-19 patients' clinical data to confirm the metric.
- Collaborated with Wuhan Union Hospital and the CoronaVac team to analyze vaccination data and used immune efficacy to mechanistically predict vaccine protection rate with higher accuracy than previous works.
- Manuscript currently under review at *Nature Communications*.

Nonlinear Dynamics in Robust Yeast Cell Cycle

Peking University

Advisor: Prof. Fangting Li, School of Physics, Center for Quantitative Biology

Oct 2018 – June 2021

- Investigated all possible three-node network topologies and kinetic parameters for robust cell cycle.
- Discovered a -1/2 scaling law between S phase duration and key cyclin Clb5 caused by critical slowing down, which renders robustness to cell cycle.
- Constructed an experimental yeast strain allowing for dynamic control over Clb5 expression and visualization and timing of spindle organization.

TECHNICAL SKILLS

Programming

• Python, MATLAB, HTML, MySQL, Flask

Experiments

Molecular biology experiments; Microbe growth and conjugation experiments.

TEACHING AND OUTREACH EXPERIENCE

 Biophysics modeling of cell cycle. Biophysics Seminar, Peking University Reverse engineer the dynamics in biology. Physics Frontiers Seminar Series, Peking University 	Aug 2020 May 2019		
SELECTED AWARDS AND HONORS			
Honorable Mention Award in ICM Contest in Modeling	2021		

•	Honorable Mention Award in ICM Contest in Modeling	2021
•	Outstanding Award, Challenge Cup academic contest, Peking University	2020
•	Beijing Innovation Research Training Fellowship	2020
•	Wanglaoji Overseas Exchange Scholarship	2019
•	Outstanding Research Award, Peking University	2018