Song Feng

冯松

What I cannot create, I do not understand. — Richard P. Feynman

Positions

- 2019 now Scientist, Biological Sciences Division, Pacific Northwest National Laboratory
- 2016 2019 Postdoc, Center for Nonlinear Studies, Los Alamos National Laboratory
- 2008 2009 Research intern, CAS-MPG Partner Institute for Computational Biology

Education

- Ph.D. University of Warwick, Systems Biology, Mentor: Orkun Soyer, 2013-2016.
- M.Sc. King Abdullah University of Science and Technology, Biosciences, 2009-2011.
- B.Sc. Zhejiang University, Biotechnology, 2005-2009.

Research Interests

General Complex (bio)systems, evolution, control, synthetic biology.

Current Biological systems: biological networks, dynamics, control, microbiome, metabolism Computational methods: modeling, inference, optimization, machine learning Social systems: opinion dynamics, social contagion

Talks

Invited

- 05/2018 Department of Mechanical engineering, University of New Mexico, United States.
- 05/2015 Evolution and optimality in cellular systems workshop, FachInstitut für Theoretische Biologie, Humboldt-Universität zu Berlin, Germany.

Contributed

09/2016 International Conference on Systems Biology (ICSB), Barcelona, Spain.

Selected Publication

* Equal contribution; † Corresponding.

Peer-reviewed

- 9 Suderman R, Mitra ED, Lin YT, Erickson KE, Feng S, Hlavacek HS[†], (2019) Generalizing Gillespie's direct method to enable network-free simulations, *Bulletin of Mathematical Biology* 81 (8), 2822-2848. doi:10.1007/s11538-018-0418-2
- 8 Lin YT*, Feng S*, Hlavacek WS[†], (2019) Acceleration of stochastic simulations of biochemical systems via heterogeneous and adaptive scaling of reaction rates, *The Journal of Chemical Physics* 150 (24), 244101. doi:10.1063/1.5096774

- 7 Feng S, Soyer OS[†], (2019) In silico evolution of signaling networks using rule-based models: bistable response dynamics, Methods Mol Biol. 1945:315-339. doi:10.1007/978-1-4939-9102-0_15
- 6 Shirin A*, Klickstein IS*, Feng S*, Lin YT*, Hlavacek WS[†], Sorrentino F[†], (2019) Prediction of Optimal Drug Schedules for Controlling Autophagy, *Sci. Rep.* 9(1):1428. doi:10.1038/s41598-019-38763-9
- 5 Jiang X, Zerfa C, Feng S, Eichmann R, Asally M, Shafer P^{\dagger} , Soyer OS^{\dagger} (2018) Impact of spatial organization on a novel auxotrophic interaction among soil microbes, *ISME J.* 12(1):1443-1456. doi:10.1038/s41396-018-0095-z
- 4 Cavaliere M, Feng S, Soyer OS, Jimenez JI[†] (2017) Cooperation in microbial communities and their biotechnological applications, *Environ. Microbiol.* 19(8):2949-2963. doi:10.1111/1462-2920.13767
- 3 Feng S*, Sáez M*, Wiuf C, Feliu E†, Soyer OS† (2016) Core signalling motif displaying multistability through multi-state enzymes. J. R. Soc. Interface 13 (123), 20160524. doi:10.1098/rsif.2016.0524
- 2 Feng S, Ollivier JF, Soyer OS[†] (2016) Enzyme sequestration as a tuning point in controlling response dynamics of signalling networks. PLoS Comput. Biol. 12(5): e1004918. doi:10.1371/journal.pcbi.1004918
- 1 Feng S*, Ollivier JF*, Swain PS[†], Soyer OS[†] (2015) BioJazz: in silico evolution of cellular networks with unbounded complexity using rule-based modeling. Nucleic Acids Res., pp. 1-15. ISSN 0305-1048 doi:10.1093/nar/gkv595.