



Universiteit  
Leiden

The Netherlands

## Applications of multisource data-based dynamic modeling to cell-cell signaling and infectious disease spreading

Chen, D.

### Citation

Chen, D. (2024, January 9). *Applications of multisource data-based dynamic modeling to cell-cell signaling and infectious disease spreading*. Retrieved from <https://hdl.handle.net/1887/3677323>

Version: Publisher's Version

License: [Licence agreement concerning inclusion of doctoral thesis in the Institutional Repository of the University of Leiden](#)

Downloaded from: <https://hdl.handle.net/1887/3677323>

**Note:** To cite this publication please use the final published version (if applicable).

# Publication list

## Work presented in this thesis

1. **Daipeng Chen**, Yanni Xiao and Sanyi Tang. “Air quality index induced non-smooth system for respiratory infection.” *Journal of Theoretical Biology* 460 (2019): 160-169.
2. **Daipeng Chen**, Yuyi Xue and Yanni Xiao. “Determining travel fluxes in epidemic areas.” *PLoS computational biology* 17.10 (2021): e1009473.
3. **Daipeng Chen**, Zary Forghany, Xinxin Liu, Haijiang Wang, Roeland Merks and David Baker. “A new model of Notch signalling: Control of Notch receptor cis-inhibition via Notch ligand dimers.” *PLoS computational biology* 19.1 (2023): e1010169.
4. **Daipeng Chen**, Xiaodan Sun and Robert A. Cheke. “Inferring a causal relationship between environmental factors and respiratory infections using convergent cross-mapping.” *Entropy* 25.5 (2023): 807.

## Other work

1. Xue, Yuyi, **Daipeng Chen**, Stacey R. Smith, Xiaoe Ruan, and Sanyi Tang. “Coupling the Within-Host Process and Between-Host Transmission of COVID-19 Suggests Vaccination and School Closures are Critical.” *Bulletin of Mathematical Biology* 85.1 (2023): 6.
2. Shixiong Cheng, Chris Jacobs, Elisa Pérez, **Daipeng Chen**, Joep Sanden, Kevin Bretscher, Femke Verweij, Jelle Bosman, Amke Hackmann, Roeland Merks,

## Publication list

---

Joost Heuvel and Maurijn Zee. “A life-history allele of large effect shortens developmental time in a wild insect population.” *Nature Ecology and Evolution* (2023): 1-13.

# Acknowledgements

During my PhD, I spent a bittersweet scientific research time, met supervisors who benefited my life, and made friends with whom I could share my joys and frustrations. I would like to thank everyone who helped me and supported me in the past years of my graduate life.

First of all, I would like to thank my supervisors, Prof. Roeland Merks from Leiden University, Prof. Yanni Xiao from Xi'an Jiaotong University and Dr. David Baker from Leiden University Medical Center. Thank you all for the enjoyable scientific discussions, for giving me the freedom to establish research questions, and for providing me timely guidance when I got lost. Prof. Xiao, thank you for introducing me and encouraging me to the wonderful scientific study. Prof. Merks, thank you for your support and your contagious enthusiasm for scientific research. Dr. Baker, thank you for your insightful suggestions on biological problem and for always making time for me. I would also like to thank my student Jeroen Nouwens whose work helped me a lot to finish the Chapter 4 of my thesis.

Next, I would like to thank Suzuki san, the Silk family and my colleagues: Erika Tsingos, Alice Peng, Koen Keijzer, Martijn de Jong, Tessa Vergroesen, David Versluis and Daphne Nesenberend from Leiden University. During the more than two years of living in the Netherlands, you provided me with help and companionship, which made me less lonely in a foreign country. In addition, I would like to thank Haijiang Wang, Xinxin Liu and Zary Forghany from Leiden University Medical Center for their contribution to my modeling work on Notch signaling. I would also like to acknowledge my collaborators Prof. Sanyi Tang, Prof. Robert Cheke and Dr. Xiaodan Sun, for helping me complete my research stories.

Thirdly, I would like thank the group members in Xi'an Jiaotong University. I want to thank Prof. Jian Zu, Dr. Mingwang Shen, Dr. Biao Tang, Dr. Pengfei Song and many others for their suggestion and comments on my academic research. Moreover,

## **Acknowledgements**

---

I have to give thanks to my friends, who shared the happiness as well as helped me out of the hard time. Yunhu Zhang, Qian Li, Fan Xia, Miaolei Li, Jun Shu, Ruikang Pang, Zhixiong Ying and Shixiong Cheng, I'll never forget the time we played and studied together.

I must express my heartfelt gratitude to my parents and my brother for their unwavering support and love over the past several years. Your understanding, encouragement, and support have been the cornerstone of my achievements, and without you, completing this thesis would not have been possible. I am deeply grateful for the collaboration and love of my girlfriend, Yuyi Xue, whose presence always provides a sense of relaxation and hopefulness.

# Curriculum Vitae

Daipeng Chen was born and brought up in Ankang, China, and attended high school at his hometown. He went to study mathematics at Northwest University, Xi'an, China in 2012 and received his BSc degree in the summers of 2016. In the fall of 2016, he started his research as a master student at the School for Mathematics and Statistics of Xi'an Jiaotong University (XJTU). During this period, he went to Shanghai for three months to learn biology at the Shanghai Institute of Biochemistry and Cell Biology, Chinese Academy of Sciences. In 2018 he began his doctoral research on modeling the transmission of infectious disease, under the supervision of Prof. Yanni Xiao at XJTU. In 2020, sponsored by the China Scholarship Council, he joined the double-PhD program of XJTU and Leiden University (LU), the Netherlands where he was supervised by Prof. Roeland Merks at the Mathematical Institute of Leiden University. He is also co-supervised by Dr. David Baker of Leiden University Medical Center. At this stage, his research focuses on developing mathematical models to investigate the mechanisms of Notch signaling and its role in angiogenesis. He is also active in teaching: at LU he has supervised a Bachelor student on the topic of "Mathematical modeling of Notch ligands heterodimerization during angiogenesis". As a teaching assistant, He has taught "Linear Algebra (XJTU)", "Systems Biology (LU)", "Physics and Chemistry for biodynamic modelling (LU)" and "Elementary and Practical Mathematics for non-mathematicians (LU)".