# **Curriculum Vitae**

Name Guangyuan(Frank) Li

Position Postdoctoral Fellow, NYU Grossman School of Medicine

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Profile <u>Blog</u>, <u>GitHub</u>, <u>Linkedin</u>

### Skill Sets

- 1. Deep Learning (Pytorch, Tensorflow, Keras) and probabilistic modeling (pyMC, pyro)
- 2. NGS analysis: DNA, RNA, (immuno)proteomics, spatial, imaging, TCR, single-cell
- 3. Target Discovery, Neoantigen pipeline, mRNA optimization
- 4. Gene/Splicing Regulatory Network
- 5. Web development (HTML, CSS, JavaScript, Flask, Dash, MySQL)
- 6. Python, Linux, R, Matlab, C, Data Visualization
- 7. Docker, Singularity, cloud computing (AWS, GCP) [Example]
- 8. Code Documentation [Example]
- 9. Quick and continual Learner.

### Education

08/2019 - 08/2023 PhD student, Division of Biomedical Informatics
Cincinnati Children's Hospital Medical Center, United States

09/2018 - 04/2019 Exchange Student, Biodesign Institute
Arizona State University, United States

09/2015 - 06/2019 Bachelor of Science, Division of Life Science Wuhan University, China

### **Working Experience**

09/2023 - Present Postdoctoral Fellow, New York University, New York, NY, United States

 Building a comprehensive target discovery pipeline for developing peptide-centric CAR-T therapy (PC-CAR)

05/2022 - 08/2022 Bioinformatics Intern, Sanofi, Cambridge, MA, United States

- Evaluating spatial deconvolution methods on 10x Visium data to guide the drug target selection and validation
- Developing standardized spatial analysis framework on AWS server to support bench scientists analysis

#### **Patent Applications**

 Nathan Salomonis, <u>Guangyuan Li</u>. "Identification of multiple targets for immunotherapy in melanoma using splicing-derived neoantigens". *PCT Patent Application No.* PCT/US2024/44819

- Mark Yarmarkovich, Zachary Harpaz, <u>Guangyuan Li</u>, Michele Palamenghi. "Systems, methods and computer-accessible medium for identifying target pairs for CAR-T therapy." U.S. Patent Application 18/940,364
- 3. Mark Yarmarkovich, <u>Guangyuan Li</u>. "Methods and compositions for use of tumor specific antigens in adoptive immunotherapy." *A PCT Patent Application has been filed and is pending*

## **First-Author Publications**

- Guangyuan Li\*, Omar Uriel Guzman-Bringas, Aman Sharma, Zachary Harpaz, Ethan Tardio, Aswin Natarajan, Xinya Liu, Darren Taylor, Rachel Yamin, Maxence Dellacherie, Michelle Krogsgaard, Conrad Russell Cruz, Mark Yarmarkovich "A pan-cancer atlas of T cell targets", *BioRxiv*. 2025. https://www.biorxiv.org/content/10.1101/2025.01.22.634237v1
- 2. **Guangyuan Li\***, Shweta Mahajan\*, Siyuan Ma, Erin D. Jeffery, Xuan Zhang, Anukana Bhattacharjee, Meenakshi Venkatashubramanian et al. "Splicing Neoantigen Discovery with SNAF Reveals Shared Targets for Cancer Immunotherapy." *Science Translational Medicine*. 2024. https://doi.org/10.1126/scitranslmed.ade2886.
- 3. <u>Guangyuan Li\*,</u> Daniel Schnell, Anukana Bhattacharjee, Mark Yarmarkovich, Nathan Salomonis. "Quantifying tumor specificity using Bayesian probabilistic modeling for drug and immunotherapeutic target discovery." *Cell Reports Methods*. 2024. <a href="https://doi.org/10.1016/j.crmeth.2024.100900">https://doi.org/10.1016/j.crmeth.2024.100900</a>
- Guangyuan Li\*, Balaji Iyer, V.B. Surya Prasath, YiZhao Ni, Nathan Salomonis. "DeepImmuno: Deep Learning-Empowered Prediction and Generation of Immunogenic Peptides for T-Cell Immunity." *Briefings in Bioinformatics*. 2021. <a href="https://doi.org/10.1093/bib/bbab160">https://doi.org/10.1093/bib/bbab160</a>.
- Guangyuan Li\*, Baobao Song, Harinder Singh, V.B. Surya Prasath, H. Leighton Grimes, Nathan Salomonis. "Decision level integration of unimodal and multimodal single cell data with scTriangulate" *Nature Communications*. 2023. <a href="https://doi.org/10.1038/s41467-023-36016-y">https://doi.org/10.1038/s41467-023-36016-y</a>
- Guangyuan Li\*, Nathan Salomonis. "RNA Isoforms as Broad Targets for Cancer Immunotherapy." DNA and Cell Biology. 2024. https://www.liebertpub.com/doi/10.1089/dna.2024.0108
- Guangyuan Li\*, Amir Bayegan, Joon Sang Lee, Donald Jackson, Jack Pollard. "Evaluating diverse deconvolution methods for tumor spatial transcriptomic datasets." Journal for ImmunoTherapy of Cancer. 2022. https://doi.org/10.1136/jitc-2022-SITC2022.0926

# **Collaborative Publications**

8. Minzhe Guo, Michael P. Morley, Cheng Jiang, Yixin Wu, <u>Guangyuan Li</u>, Yina Du, Shuyang Zhao et al. "Guided Construction of Single Cell Reference for Human and Mouse Lung." *Nature Communications*. 2023. https://www.nature.com/articles/s41467-023-40173-5

- 9. Kang Jin, Daniel Schnell, Guangyuan Li, Nathan Salomonis, V.B. Surya Prasath, Rhonda Szczesniak, Bruce J. Aronow. "CellDrift: Inferring Perturbation Responses in Temporally-Sampled Single Cell Data." Briefing in Bioinformatics. 2022. https://doi.org/10.1093/bib/bbac324
- 10. Zhang Xuan, Baobao Song, Maximillian J. Carlino, Guangyuan Li, Kyle Ferchen, Mi Chen, Evrett N. Thompson, et al. "An Immunophenotype-Coupled Transcriptomic Atlas of Human Hematopoietic Progenitors." Nature Immunology. 2024. https://www.nature.com/articles/s41590-024-01782-4
- 11. L. Lambourne, K. Mattioli, C. Santoso, G. Sheynkman, S. Inukai, B. Kaundal, A. Berenson, K. Spirohn-Fitzgerald, A. Bhattacharjee, E. Rothman, S. Shrestha, F. Laval, Z. Yang, D. Bisht, J. A. Sewell, G. Li, A. Prasad, S. Phanor, R. Lane, D. M. Campbell, T. Hunt, D. Balcha, M. Gebbia, J.-C. Twizere, T. Hao, A. Frankish, J. A. Riback, N. Salomonis, M. A. Calderwood, D. E. Hill, N. Sahni, M. Vidal, M. L. Bulyk, J. I. Fuxman Bass "Widespread variation in molecular interactions and regulatory properties among transcription factor isoforms". bioRxiv, 2024.
  - https://www.biorxiv.org/content/10.1101/2024.03.12.584681v2