

Minghao JIANG | 姜明昊

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EDUCATION

- **Ph.D. in Bioinformatics at The University of Hong Kong**, Hong Kong, Hong Kong SAR Sep-2023 — now
 - Department: School of Biomedical Sciences, Li Ka Shing Faculty of Medicine
 - Supervisor: Professor Jason Wing Hon WONG
- **M.Sc. in Biology at Shanghai Jiao Tong University**, Shanghai, China Sep-2019 — Jun-2022
 - Department: Shanghai Institute of Hematology, State Key Laboratory of Medical Genomics
 - Supervisor: Professor Lin CHENG
 - GPA: 3.90/4.0
- **B.Sc. in Bioinformatics at Harbin Medical University**, Harbin, Heilongjiang Province, China Sep-2014 — Jul-2019
 - College: College of Bioinformatics Science and Technology
 - Average score: 81.41/100 (five-year program)

RESEARCH INTERESTS

Bioinformatics | Computational biology | Genome organization | RNA biology

RESEARCH EXPERIENCE

1. Identified dysregulated sub-pathways in cancers by integrating gene expressions, CNVs, and DNA methylations.
2. Identified cell-cell communications associated with prognosis of patients with B-cell acute lymphoblastic leukemia (B-ALL).
3. Revealed TCF12 acted as a co-factor of IGH::DUX4 to enhance the transcativation via a positive feedback in B-ALL.
4. Benchmarked 21 differential splicing tools for RNA-seq analysis at the event level to propose a tool-selection protocol for users.
5. Predicted the spreading of the COVID-19 epidemic in China using a novel time-delay differential equation model.
6. Performed analysis of RNA-seq, ChIP-seq, scRNA-seq, and CyTOF data.

PUBLICATIONS ¹

Xue Q[†], Zhang M[†]✉, Mo Y[†], Jiao B[†], Liu X[†], Jiang M, Zhou Y, Tan Y, Li H, Zhang J, Zhang Q, Li Y, Li J, Ma X, Jing DH, Mi JQ, Wang J, Chen Z, Shen SH[✉], Chen SJ[✉]. **Molecular mechanisms of unique therapeutic potential of CUDC-907 for MEF2D fusion-driven BCP-ALL.** *Signal Transduct Target Ther.* 2025 Jul 23;10(1):230. [10.1038/s41392-025-02310-y](https://doi.org/10.1038/s41392-025-02310-y)

Jiang M, Jing D, Wong JWH[✉]. **gghic: A Versatile R Package for Exploring and Visualizing 3D Genome Organization.** *arXiv.* 2024 Dec. [10.48550/arXiv.2412.03005](https://arxiv.org/abs/10.48550/arXiv.2412.03005)

Ye J[†], Wang F[†], Zhou Z[†], Jiang M[†], You Y, Su Y, Hu Q, Chi H, Liu H, Cheng X, Jia J, Shi H, Yang C, Huang J[⊗], Teng J[⊗], Hu Q[⊗], Sun Y[⊗]. **Itaconate promoted monocyte migration by increasing chemokine CCL12 expression in autoinflammatory cytokine storm.** *Clin Immunol.* (Under review)

Li Z[†], Jiang M[†], Wang J[†], Zhuo Z[†], Zhang S, Tan Y, Hu W[⊗], Zhang H[⊗], Meng G[⊗]. **Transcription factor 12-mediated self-feedback regulatory mechanism is required in *DUX4* fusion leukaemia.** *Clin Transl Med.* 2023 Dec;13(12):e1514. [10.1002/ctm2.1514](https://doi.org/10.1002/ctm2.1514).

Yin H[†], Wang J[†], Tan Y[†], Jiang M[†], Zhang H[⊗], Meng G[⊗]. **Transcription factor abnormalities in B-ALL leukemogenesis and treatment.**[#] *Trends Cancer.* 2023 Oct;9(10):855-870. [10.1016/j.trecan.2023.06.004](https://doi.org/10.1016/j.trecan.2023.06.004).

Wu W[†], Tan Y[†], Yin H[†], Jiang M[†], Jiang Y, Ma X, Yin T, Li Y[⊗], Zhang H[⊗], Cai X[⊗], Meng G[⊗]. **Phase separation is required for PML nuclear body biogenesis and function.** *FASEB J.* 2023 Jun;37(6):e22986. [10.1096/fj.202300216R](https://doi.org/10.1096/fj.202300216R).

Jiang M^{†⊗}, Zhang S[†], Yin H[†], Zhuo Z[†], Meng G[⊗]. **A comprehensive benchmarking of differential alternative splicing tools at the event level.** *Brief Bioinform.* 2023 May 19;24(3):bbad121. [10.1093/bib/bbad121](https://doi.org/10.1093/bib/bbad121).

Lu G[†], Ling Y[†], Jiang M[†], Tan Y[†], Wei D[†], Jiang L, Yu S, Jiang F, Wang S, Dai Y, Wang J, Wu G, Zhang X, Meng G[⊗], Wang S[⊗], Liu F[⊗], Fan X[⊗], Chen S[⊗]. **Primary assessment of the diversity of Omicron sublineages and the epidemiologic features of autumn/winter 2022 COVID-19 wave in Chinese mainland.** *Front Med.* 2023 Aug;17(4):758-767. [10.1007/s11684-022-0981-7](https://doi.org/10.1007/s11684-022-0981-7).

Jiang M[†], Yin H[†], Zhang S[†], Meng G[⊗], Wu G[⊗]. **Mathematical appraisal of SARS-CoV-2 Omicron epidemic outbreak in unprecedented Shanghai lockdown.** *Front Med (Lausanne).* 2022 Nov 8;9:1021560. [10.3389/fmed.2022.1021560](https://doi.org/10.3389/fmed.2022.1021560).

Zhang M[†], Zhang H[†], Li Z[†], Bai L[†], Wang Q, Li J, Jiang M, Xue Q, Cheng N, Zhang W, Mao D, Chen Z, Huang J, Meng G[⊗], Chen Z[⊗], Chen SJ[⊗]. **Functional, structural, and molecular characterizations of the leukemogenic driver MEF2D-HNRNPUL1 fusion.** *Blood.* 2022 Sep 22;140(12):1390-1407. [10.1182/blood.2022016241](https://doi.org/10.1182/blood.2022016241).

Zhang H[†], Cheng N[†], Li Z[†], Bai L[†], Fang C, Li Y, Zhang W, Dong X, Jiang M, Liang Y, Zhang S, Mi J, Zhu J, Zhang Y, Chen SJ, Zhao Y, Weng XQ, Hu W[⊗], Chen Z[⊗], Huang J[⊗], Meng G[⊗]. **DNA crosslinking and recombination-activating genes 1/2 (RAG1/2) are required for oncogenic splicing in acute lymphoblastic leukemia.** *Cancer Commun (Lond).* 2021 Nov;41(11):1116-1136. [10.1002/cac2.12234](https://doi.org/10.1002/cac2.12234).

Wu L[†], Jiang M[†], Yu P, Li J, Ouyang W, Feng C, Zhao WL[⊗], Dai Y[⊗], Huang J[⊗]. **Single-Cell Transcriptome Analysis Identifies Ligand-Receptor Pairs Associated With BCP-ALL Prognosis.** *Front Oncol.* 2021 Mar 10;11:639013. [10.3389/fonc.2021.639013](https://doi.org/10.3389/fonc.2021.639013).

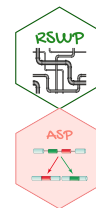
Han J^{†⊗}, Liu S[†], Jiang Y[†], Xu C[†], Zheng B, Jiang M, Yang H, Su F, Li C[⊗], Zhang Y[⊗]. **Inference of patient-specific subpathway activities reveals a functional signature associated with the prognosis of patients with breast cancer.** *J Cell Mol Med.* 2018 Sep;22(9):4304-4316. [10.1111/jcmm.13720](https://doi.org/10.1111/jcmm.13720).

SOFTWARE & TOOLS

- [gghic](#)
A versatile R package for exploring and visualizing 3D genome organization



- [smk_sv](#)
A **s**nakemake pipeline to call **s**tructural **v**ariants from tumor-only ONT data
- [RSWP](#)
RNA sequencing workflow by Python
- [ASP](#)
An **a**lternative **s**plicing **p**ackage in R
- Membership in the [openbioX](#) community



AWARDS

- | | |
|---|------|
| • First-class scholarship, Shanghai Jiao Tong University School of Medicine | 2021 |
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| • Outstanding graduate, Harbin Medical University | 2019 |
| • Outstanding volunteer of The Belt and Road China-Russia Medical University Alliance | 2018 |
| • Excellent student leader, Harbin Medical University | 2016 |

OTHER ROLES

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| • Director of Sports Department, Student Union, Shanghai Jiao Tong University School of Medicine | 2019 — 2020 |
| • Director of Publicity Department, Student Union, Harbin Medical University | 2014 — 2016 |

SKILLS

- Proficient in R, Python, Shell, Linux, Slurm (Simple Linux Utility for Resource Management), and Adobe Illustrator
- Familiar with Perl, Java, and Adobe Photoshop