

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 transactivation 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

Qing Xue[†], Ming Zhang[†], Yixiao Mo[†], Bo Jiao[†], Xuan Liu[†], **Minghao Jiang**, Yu Zhou, Yun Tan, Huimin Li, Jianming Zhang, Qianqian Zhang, Yunqi Li, Jianfeng Li, Xiaofang Ma, Duo-Hui Jing, Jian-Qing Mi, Jin Wang, Zhu Chen, Shu-Hong Shen[✉], Sai-Juan Chen[✉]. "Molecular Mechanisms of Unique Therapeutic Potential of CUDC-907 for *MEF2D* Fusion-driven BCP-ALL" *Signal Transduct Target Ther.* (Accepted on June 23rd, 2025)

Minghao Jiang, Duohui Jing, Jason W.H. Wong[✉]. "gghic: A Versatile R Package for Exploring and Visualizing 3D Genome Organization" 2024 Dec. *arXiv* doi: [10.48550/arXiv.2412.03005](https://doi.org/10.48550/arXiv.2412.03005)

Junna Ye[†], Fan Wang[†], Zhuochao Zhou[†], **Minghao Jiang**[†], Yijun You, Yutong Su, Qiongyi Hu, Huihui Chi, Honglei Liu, Xiaobing Cheng, Jinchao Jia, Hui Shi, Chengde Yang, Jinyan Huang[✉], Jialin Teng[✉], Qiongyi Hu[✉], Yue Sun[✉]. "Itaconate promoted monocyte migration by increasing chemokine CCL12 expression in autoinflammatory cytokine storm." *Clin Immunol.* (Under review)

Zhihui Li[†], **Minghao Jiang**[†], Junfei Wang[†], Zhiyi Zhuo[†], Shiyan Zhang, Yangxia Tan, Weiguo Hu[✉], Hao Zhang[✉], Guoyu Meng[✉]. "Transcription factor 12-mediated self-feedback regulatory mechanism is required in *DUX4* fusion leukaemia." *Clin Transl Med.* 2023 Dec;13(12):e1514. doi: [10.1002/ctm2.1514](https://doi.org/10.1002/ctm2.1514).

Hongxin Yin[†], Junfei Wang[†], Yangxia Tan[†], **Minghao Jiang**[†], Hao Zhang[✉], Guoyu Meng[✉]. "Transcription factor abnormalities in B-ALL leukemogenesis and treatment."[#] *Trends Cancer.* 2023 Oct;9(10):855-870. doi:[10.1016/j.trecan.2023.06.004](https://doi.org/10.1016/j.trecan.2023.06.004).

Wenyu Wu[†], Yangxia Tan[†], Hongxin Yin[†], **Minghao Jiang**[†], Yanling Jiang, Xiaodan Ma, Tong Yin, Yuwen Li[✉], Hao Zhang[✉], Xun Cai[✉], Guoyu Meng[✉]. "Phase separation is required for PML nuclear body biogenesis and function." *FASEB J.* 2023 Jun;37(6):e22986. doi: [10.1096/fj.202300216R](https://doi.org/10.1096/fj.202300216R).

Minghao Jiang^{†, ✉}, Shiyan Zhang[†], Hongxin Yin[†], Zhiyi Zhuo[†], Guoyu Meng[✉]. "A comprehensive benchmarking of differential alternative splicing tools at the event level." *Brief Bioinform.* 2023 Apr 5;bbad121. doi: [10.1093/bib/bbad121](https://doi.org/10.1093/bib/bbad121).

Gang Lu[†], Yun Ling[†], **Minghao Jiang**[†], Yun Tan[†], Dong Wei[†], Lu Jiang, Shuting Yu, Fangying Jiang, Shuai Wang, Yao Dai, Jinzeng Wang, Geng Wu, Xinxin Zhang, Guoyu Meng[✉], Shengyue Wang[✉], Feng Liu[✉], Xiaohong Fan[✉], Saijuan Chen[✉]. "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 Mar 31;1-10. doi: [10.1007/s11684-022-0981-7](https://doi.org/10.1007/s11684-022-0981-7).

Minghao Jiang[†], Hongxin Yin[†], Shiyan Zhang[†], Guoyu Meng[✉], Geng Wu[✉]. "Mathematical appraisal of SARS-CoV-2 Omicron epidemic outbreak in unprecedented Shanghai lockdown." *Front Med (Lausanne).* 2022 Nov 8;9:1021560. doi: [10.3389/fmed.2022.1021560](https://doi.org/10.3389/fmed.2022.1021560).

Ming Zhang[†], Hao Zhang[†], Zhihui Li[†], Ling Bai[†], Qianqian Wang, Jianfeng Li, **Minghao Jiang**, Qing Xue, Nuo Cheng, Weina Zhang, Dongdong Mao, Zhiming Chen, Jinyan Huang, Guoyu Meng[✉], Zhu Chen[✉], Sai-Juan Chen[✉] "Functional, Structural and Molecular Characterizations of Leukemogenic Driver Mef2d-Hnrnpul1 Fusion." *Blood.* 2022 Sep 22;140(12):1390-1407. doi: [10.1182/blood.2022016241](https://doi.org/10.1182/blood.2022016241).

Hao Zhang[†], Nuo Cheng[†], Zhihui Li[†], Ling Bai[†], Chengli Fang, Yuwen Li, Weina Zhang, Xue Dong, **Minghao Jiang**, Yang Liang, Sujiang Zhang, Jianqing Mi, Jiang Zhu, Yu Zhang, Sai-Juan Chen, Yajie Zhao, Xiang-Qin Weng, Weiguo Hu[✉], Zhu Chen[✉], Jinyan Huang[✉], Guoyu Meng[✉]. "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. doi: [10.1002/cac2.12234](https://doi.org/10.1002/cac2.12234).

Liang Wu[†], **Minghao Jiang**[†], Ping Yu, Jianfeng Li, Wen Ouyang, Chong Feng, Wei Li Zhao[✉], Yuting Dai[✉], Jinyan Huang[✉]. "Single-Cell Transcriptome Analysis Identifies Ligand-Receptor Pairs Associated with BCP-ALL Prognosis." *Front Oncol.* 2021 Mar 10;11:639013. doi: [10.3389/fonc.2021.639013](https://doi.org/10.3389/fonc.2021.639013).

Junwei Han^{†, ✉}, Siyao Liu[†], Ying Jiang[†], Chaohan Xu[†], Baotong Zheng, **Minghao Jiang**, Haixiu Yang, Fei Su, Chunquan Li[✉], Yunpeng Zhang[✉]. "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. doi: [10.1111/jcmm.13720](https://doi.org/10.1111/jcmm.13720).

PUBLISHED SOFTWARE

- [gghic](#)
A versatile R package for exploring and visualizing 3D genome organization
- [smk_sv](#)
A **snakemake** pipeline to call **structural variants** from tumor-only ONT data
- [RSWP](#)
RNA sequencing workflow by **Python**
- [ASP](#)
An **alternative splicing package** 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

1. Time-ordered | [†] Co-first authors | [✉] Corresponding author(s) | [#] Review articles ↩