

# Rui Yin

yinrui2000@outlook.com | [frickyinn.github.io](https://frickyinn.github.io)

## Education

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|--|-----------------------|
| <b>The University of Hong Kong</b> <ul style="list-style-type: none"><li>• Master in Science, Computer Science</li><li>• Withdrawn</li></ul>         | Sept 2022 – Jan 2023  |
| <b>Sun Yat-sen University</b> <ul style="list-style-type: none"><li>• Bachelor of Engineering, Software Engineering</li><li>• GPA: 3.6/4.0</li></ul> | Sept 2018 – June 2022 |

## Interests

Computer Vision, Machine Learning, 3D Vision, Deep Model Compression

## Publications

### SRPose: Two-view Relative Pose Estimation With Sparse Keypoints

Rui Yin, Yulun Zhang, Zherong Pan, Jianjun Zhu, Cheng Wang, Biao Jia  
ECCV, 2024  
[project page](#)

### Identifying Spatial Domain by Adapting Transcriptomics With Histology Through Contrastive Learning

Yuansong Zeng\*, Rui Yin\*, Mai Luo, Jianing Chen, Zixiang Pan, Yutong Lu, Weijing Yu, Yuedong Yang  
Briefings in Bioinformatics, 2023

### Spatial Transcriptomics Prediction From Histology Jointly Through Transformer and Graph Neural Networks

Yuansong Zeng, Zhuoyi Wei, Weijiang Yu, Rui Yin, Yuchen Yuan, Bingling Li, Zhonghui Tang, Yutong Lu, Yuedong Yang  
Briefings in Bioinformatics, 2022

## Experience

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|---|--|
| <b>Hanglok-Tech</b><br>Research Assistant   Advisor: Biao Jia<br>Relative Camera Pose Estimation <ul style="list-style-type: none"><li>• SRPose: Two-view relative pose estimation with sparse keypoints</li></ul> Model Quantization <ul style="list-style-type: none"><li>• Proposing a new binarization function for model compression in depth estimation tasks (paper coming).</li></ul> | Shenzhen, China<br>Aug 2023 – Present    |
| <b>Laboratory of Data Discovery for Health, The University of Hong Kong</b><br>Research Assistant   | Hong Kong SAR<br>Oct 2022 – Jan 2023     |
| <b>AI4science Lab, Sun Yat-sen University</b><br>Research Intern   Advisor: Yuedong Yang <ul style="list-style-type: none"><li>• Identifying spatial domain by adapting transcriptomics with histology through contrastive learning</li><li>• Spatial transcriptomics prediction from histology jointly through transformer and graph neural networks</li></ul>                               | Guangzhou, China<br>Apr 2020 – Sept 2022 |

## Additional Experience

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| <b>NUS SOC 2021 Summer Workshop, National University of Singapore</b><br>Advisor: Tan Zi Qing | Singapore<br>May 2021 - Aug 2021        |
| <b>National Training Program of Innovation and Entrepreneurship</b><br>Advisor: Yuedong Yang  | Guangzhou, China<br>Apr 2020 - Nov 2020 |