

# 朱河勤

中国科学技术大学

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## 核心研究: AI FOR SCIENCE

- 计算生物学: 基于大模型的 RNA 结构与功能预测。主导开发结构引导的 RNA 基础语言模型 [I8]; 提出碱基对基序能量模型, 发表于 [Nature Communications](#) [I6]。
- 医学影像计算: 跨域通用型解剖关键点检测。研发 GU2Net [I1, I2], DATR [I4], UOD [I5], 等模型, 分别发表于 [MICCAI 2021](#), [BME Frontiers](#), [MICCAI 2023](#)。

## 教育经历

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|--|---------------------------|
| • 中国科学技术大学<br>博士研究生, 生物医学工程<br>◦ 导师: <a href="#">周少华教授</a> (Fellow of AIMBE, IAMBE, IEEE, MICCAI, and NAI) | 2023.09 - 2026 (预计)<br>苏州 |
| • 中国科学院, 计算技术研究所&中国科学院大学<br>硕士, 计算机应用技术<br>◦ 导师: <a href="#">周少华教授</a> (同上)                                | 2020.09 - 2023.06<br>北京   |
| • 中国科学技术大学<br>本科, 计算机科学与技术<br>◦ 华夏计算机科学与技术英才班  | 2016.09 - 2020.06<br>合肥   |

## 荣誉获奖

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|-----------------------|-----------|
| • 苏州工业园区奖学金, 中国科学技术大学 | 2025      |
| • 一等学业奖学金, 中国科学技术大学   | 2024-2025 |
| • 一等学业奖学金, 计算所&国科大    | 2020-2023 |
| • 三好学生, 计算所&国科大       | 2023      |
| • 优秀学生奖, 中国科学技术大学     | 2018-2019 |
| • 化研所英才奖, 中国科学技术大学    | 2017      |

## 项目实习

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|---|-------------------------|
| • 腾讯天衍实验室<br>研究实习生<br>◦ 使用深度图监督学习进行图像显著性检测, 开发了DFTR 多模态融合模型 [I3]  | 2021.07 - 2021.11<br>深圳 |
| • 课程资源整理<br><a href="#">GitHub</a> 开源项目<br>◦ 主导整理了计算机相关的学习资源, 目前获得 <a href="#">15.7K 星标</a> 。 <a href="https://github.com/USTC-Resource/USTC-Course">https://github.com/USTC-Resource/USTC-Course</a> | 2019-现在                 |

## 学术活动

- 学术审稿: MICCAI (医学影像顶会, CCF B 类)、TCSVT (IEEE 期刊, IF=5.1) : 负责 AI 在医学影像, 生物信息领域的论文评审。
- 志愿活动: 医学影像计算独墅湖会议志愿者 (2023), 医学增强现实夏季学期 (2024) : 协助组织 AI 医学影像专题论坛, 促进产学研合作。
- 助教: 电子信息开放实践: 中国科学技术大学 2023 年秋季学期。

## 技术技能

- 大模型与深度学习: PyTorch 框架, 语言模型, 扩散模型, 多模态融合, LoRA。
- 科学计算工具: 生物计算 (AlphaFold3, Protenix, PyMol), Pandas, Numpy.
- 编程语言与工具: Python, C, C++, Git, bash, VIM。

Selected publications, # denotes co-first author and \* denotes co-corresponding author. For full list, please refer to [Google Scholar](#).

### Representative Papers

- [I8] **Heqin Zhu**, RuiFeng Li, Feng Zhang, Fenghe Tang, Tong Ye, Xin Li, Yunjie Gu, Peng Xiong\*, and S. Kevin Zhou\*. "A fully open structure-guided RNA foundation model for robust structural and functional inference." ([Submitted](#)). [[bioRxiv](#); [Code](#)]
- [I7] Feng Zhang#, **Heqin Zhu#**, Jie Hu, Jiayin Gao, Ke Chen, S. Kevin Zhou\*, and Peng Xiong\*. "IRESeek: Structure-informed deep learning method for accurate identification of internal ribosome entry sites in circular RNAs." ([Submitted](#)).
- [I6] **Heqin Zhu**, Fenghe Tang, Quan Quan, Ke Chen, Peng Xiong\*, and S. Kevin Zhou\*. "Deep generalizable prediction of RNA secondary structure via base pair motif energy." *Nature Communications* 16, (2025): 5856. ([Nat. Commun., 2025](#)). [[Paper](#); [Code](#)]
- [I5] **Heqin Zhu**, Quan Quan, Qingsong Yao, Zaiyi Liu, and S. Kevin Zhou. "Uod: Universal one-shot detection of anatomical landmarks." In *International Conference on Medical Image Computing and Computer-Assisted Intervention*, pp. 24-34. Cham: Springer Nature Switzerland, 2023. ([MICCAI 2023](#)). [[Paper](#); [Code](#)]
- [I4] **Heqin Zhu**, Qingsong Yao, and S. Kevin Zhou. "Datr: Domain-adaptive transformer for multi-domain landmark detection." *arxiv preprint arxiv:2203.06433* (2022). [[Paper](#); [Code](#)]
- [I3] **Heqin Zhu**, Xu Sun, Yuexiang Li, Kai Ma, S. Kevin Zhou\*, and Yefeng Zheng\*. "DFTR: Depth-supervised fusion transformer for salient object detection." *arxiv preprint arxiv:2203.06429* (2022). [[Paper](#); [Code](#)]
- [I2] **Heqin Zhu**, Qingsong Yao, Li Xiao, and S. Kevin Zhou. "Learning to Localize Cross-Anatomy Landmarks in X-Ray Images with a Universal Model." *BME Frontiers* 2022 (2022): 9765095. ([BMEF 2022](#)). [[Paper](#); [Code](#)]
- [I1] **Heqin Zhu**, Qingsong Yao, Li Xiao, and S. Kevin Zhou. "You only learn once: Universal anatomical landmark detection." In *Medical Image Computing and Computer Assisted Intervention*, pp. 85-95. Springer International Publishing, 2021. ([MICCAI 2021](#)). [[Paper](#); [Code](#)]

### Journal Papers

- [J4] Quan Quan#, Qingsong Yao#, **Heqin Zhu**, and S. Kevin Zhou. "IGU-Aug: Information-guided unsupervised augmentation and pixel-wise contrastive learning for medical image analysis." *IEEE Transactions on Medical Imaging* (2024). ([TMI 2024](#)).
- [J3] Quan Quan#, Qingsong Yao#, **Heqin Zhu**, Qiyuan Wang, and S. Kevin Zhou. "Which images to label for few-shot medical image analysis?." *Medical Image Analysis* 96 (2024): 103200. ([MIA 2024](#)).
- [J2] Huang Zhen#, Han Li#, Shitong Shao, **Heqin Zhu**, Huijie Hu, Zhiwei Cheng, Jianji Wang, and S. Kevin Zhou. "PELE scores: pelvic X-ray landmark detection with pelvis extraction and enhancement." *International Journal of Computer Assisted Radiology and Surgery* 19, no. 5 (2024): 939-950. ([IJCAR 2024](#)).
- [J1] Pengbo Liu, Hu Han, Yuanqi Du, **Heqin Zhu**, Yinhao Li, Feng Gu et al. "Deep learning to segment pelvic bones: large-scale CT datasets and baseline models." *International Journal of Computer Assisted Radiology and Surgery* 16 (2021): 749-756. ([IJCAR 2021](#)).

### Conference Papers

- [C4] Xinyi Wang, Zikang Xu, **Heqin Zhu**, Qingsong Yao, Yiyong Sun, and S. Kevin Zhou. "SIX-Net: Spatial-Context Information miX-up for Electrode Landmark Detection." In *International Conference on Medical Image Computing and Computer-Assisted Intervention*, pp. 338-348. Cham: Springer Nature Switzerland, 2024. ([MICCAI 2024](#)).
- [C3] Fenghe Tang, Ronghao Xu, Qingsong Yao, Xueming Fu, Quan Quan, **Heqin Zhu**, Zaiyi Liu, and S. Kevin Zhou. "Hyspark: Hybrid sparse masking for large scale medical image pre-training." In *International Conference on Medical Image Computing and Computer-Assisted Intervention*, pp. 330-340. Cham: Springer Nature Switzerland, 2024. ([MICCAI 2024](#)).
- [C2] Quan Quan, Fenghe Tang, Zikang Xu, **Heqin Zhu**, and S. Kevin Zhou. "Slide-SAM: Medical SAM Meets Sliding Window." In *Medical Imaging with Deep Learning*, pp. 1179-1195. PMLR, 2024. ([MIDL 2024](#)).
- [C1] Yuanyuan Lyu, Haofu Liao, **Heqin Zhu**, and S. Kevin Zhou. "A 3 DSegNet: anatomy-aware artifact disentanglement and segmentation network for unpaired segmentation, artifact reduction, and modality translation." In *International Conference on Information Processing in Medical Imaging*, pp. 360-372. Cham: Springer International Publishing, 2021. ([IPMI 2021](#)).