

Qiao Liu – Curriculum Vitae

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Research Statement

My research interests cover problems on statistical learning and computational biology problems. In particular, I am interested in developing machine learning algorithms, especially deep learning algorithms with applications in both general statistics and specific problems in computational biology, pharmacopathology, and health care. Related related works have been published in top journals (**PNAS, Nature Machine Intelligence, Bioinformatics**) and conferences (**ISMB, ECCB, NeurIPS, MICCAI**).

Employment

2021.06- Postdoctoral Scholar - Stanford University, Stanford, USA
Present Department of Statistics, advised by Prof. **Wing Hung Wong** (NAS member)
Developing ML algorithms for solving problems in general statistics and computational biology.

Education

2019.09- Joint Ph.D. student - Stanford University, Stanford, USA
2021.06 Department of Statistics, advised by Prof. **Wing Hung Wong** (NAS member)
Developing machine learning algorithms with applications in both general statistics and biomedical data

2016.09- Ph.D. candidate in **Control Science and Engineering** - Tsinghua University, Beijing, China
2021.06 Department of Automation, advised by tenured Assoc. Prof. **Rui Jiang**
Tsinghua National Laboratory for Informatics and Technology (TNLIST)

2015.08- Exchange Student - Lund University, Lund, Sweden
2016.01 Department of Computer Science
GPA:**5.0**/5.0

2012.09- Bachelor in **Engineering** - Beihang University, Beijing, China
2016.06 ShenYuan Honors College (Special Pilot Class, formerly a.k.a. School of Advanced Engineering)
GPA:**91.5**/100, Rank:**2**/50

Internships

2019.06- Research Intern - SenseTime Inc, Beijing, China
2019.09 AI+Healthcare, mentored by Dr. **Mu Zhou**
Developing ML algorithms for drug sensitivity prediction and novel drug discovery

2015.09- Undergraduate Intern - LUGG Lab, Lund, Sweden
2015.12 Lund University Graphics Group, advised by Assoc. Prof. **Michael Doggett**
Developing and maintaining RenderChimp graphics applications platform

Publications

(†=co-first author; *=corresponding author)

Preprints

1. Qijin Yin, Xusheng Cao, Rui Fan, **Qiao Liu**^{*}, Rui Jiang^{*} and Wanwen Zeng^{*}. DeepDrug: A general graph-based deep learning framework for drug-drug interactions and drug-target interactions prediction[J]. bioRxiv, 2022.
2. Kexin Ding, Mu Zhou, Zichen Wang, **Qiao Liu**, Corey W. Arnold, Shaoting Zhang^{*}, Dimitri N. Metaxas^{*}. Graph Convolutional Networks for Multi-modality Medical Imaging: Methods, Architectures, and Clinical Applications[J]. arXiv preprint arXiv:2202.08916, 2022.

Peer-reviewed papers

1. **Qiao Liu**, Kui Hua, Xuegong Zhang, Wing Hung Wong^{*}, Rui Jiang^{*}. DeepCAGE: incorporating transcription factors in genome-wide prediction of chromatin accessibility[J]. *Genomics, Proteomics & Bioinformatics*, 2022.
2. **Qiao Liu**, Jiase Xu, Rui Jiang^{*}, Wing Hung Wong^{*}. Density estimation with deep generative neural networks [J]. Proceedings of the National Academy of Sciences of the United States of America (*PNAS*), 2021, 118(15), e2101344118.
3. **Qiao Liu**, Shengquan Chen, Rui Jiang^{*}, Wing Hung Wong^{*}. Simultaneous deep generative modeling and clustering of single cell genomic data[J]. *Nature Machine Intelligence*, 2021, 3(6): 536-544.
4. **Qiao Liu**, Zhiqiang Hu, Rui Jiang^{*}, Mu Zhou^{*}. Cancer drug response prediction via a hybrid graph convolutional network[J]. *Bioinformatics*, 2020. 36(Supplement_2): i911-i918. Also in *Proceedings of the 19th European Conference on Computational Biology (ECCB)*, 2020. (Top bioinformatics conf, conference acceptance rate:20.2%)
5. **Qiao Liu**, Hairong Lv, Rui Jiang^{*}. hicGAN infers super resolution Hi-C data with generative adversarial networks. *Bioinformatics*, 2019, 35(14): i99-i107. Also in *Proceedings of the 27th Intelligent Systems for Molecular Biology/the 18th European Conference on Computational Biology (ISMB/ECCB)*, 2019. (Top bioinformatics conf, conference acceptance rate:18.9%)
6. **Qiao Liu**, Fei Xia, Qijin Yin, Rui Jiang^{*}. Chromatin accessibility prediction via a hybrid deep convolutional neural network[J]. *Bioinformatics*, 2017, 34(5): 732-738. (Top bioinformatics journal, JCR Q1)
7. **Qiao Liu**, Mingxin Gan, Rui Jiang^{*}. A sequence-based method to predict the impact of regulatory variants using random forest[J]. *BMC Systems Biology*, 2017, 11(2): 7. (JCR Q2) Also in *Proceedings of the 15th Asia Pacific Bioinformatics Conference (APBC)*, 2017.
8. Pengyu Chen[†], **Qiao Liu**[†], Lan Wei, Beier Zhao, Yin Jia, Hairong Lv^{*}, Xiaolu Fei^{*}. Automatically structuring on Chinese ultrasound report of cerebrovascular diseases via natural language processing[J]. *IEEE Access*, 2019, 7: 89043-89050. (Co-first author, JCR Q1)
9. Qijin Yin, **Qiao Liu**, Zhuoran Fu, Rui Jiang^{*}. scGraph: a graph neural network-based approach to automatically identify cell types[J]. *Bioinformatics*, 2022. (Top bioinformatics journal, JCR Q1)
10. Shengquan Chen, **Qiao Liu**, Xuejian Cui, Rui Jiang^{*}. OpenAnnotate: a web server to annotate the chromatin accessibility of genomic regions[J]. *Nucleic Acids Research*, 2021, 49(W1): W483-W490. (JCR Q1)
11. Feng Wang, Guoyizhe Wei, **Qiao Liu**, Jinxiang Ou, Xian Wei, Hairong Lv^{*}. Boost Neural Networks by Checkpoints [C]. Conference on Neural Information Processing Systems (*NeurIPS*), 2021, 33. (Top machine learning conf, acceptance rate:26%)
12. Chencheng Xu, **Qiao Liu**, Minlie Huang, Tao Jiang^{*}. Reinforced molecular optimization with neighborhood-controlled grammars[C]. Conference on Neural Information Processing Systems (*NeurIPS*), 2020, 33. (Top machine learning conf, acceptance rate:20.1%)
13. Kexin Ding, **Qiao Liu**, Edward Lee, Mu Zhou, Aidong Lu, Shaoting Zhang^{*}. Feature-enhanced graph networks for genetic mutational prediction using histopathological images in colon cancer[C]. International Conference on Medical Image Computing and Computer Assisted Intervention (*MICCAI*), 2020, (pp. 294-304). (Top medical AI conf)

14. Chencheng Xu, **Qiao Liu**, Jianyu Zhou, Minzhu Xie, Jianxing Feng, Tao Jiang*. Quantifying functional impacts of regulatory variants with multi-task Bayesian neural network[J]. *Bioinformatics*, 2020, 36(5): 1397-1404. (Top bioinformatics journal, JCR Q1)
15. Qingzhu Yang, **Qiao Liu**, Hairong Lv*. A Decentralized System for Medical Data Management via Blockchain [J]. *Journal of Internet Technology*, 2020, 21(5): 1335-1345.
16. Junfeng Liu, **Qiao Liu**, Qingzhu Yang*. Mstree: a multispecies coalescent approach for estimating ancestral population size and divergence time during speciation with gene flow [J]. *Genome biology and evolution*, 2020, 12(5): 715-719. (JCR Q2)
17. Shaoming Song, Hongfei Cui, Shengquan Chen, **Qiao Liu**, Rui Jiang*. EpiFIT: Functional interpretation of transcription factors based on combination of sequence and epigenetic information[J]. *Quantitative Biology*, 2019, 1-11.
18. Qijin Yin, Mengmeng Wu, **Qiao Liu**, Rui Jiang*. DeepHistone: a deep learning approach to predicting histone modifications[J]. *BMC Genomics*, 2019,20(2):193. (JCR Q2)
19. Bai Li, Mu Lin, **Qiao Liu**, Ya Li*, Changjun Zhou*. Protein folding optimization based on 3D off-lattice model via an improved artificial bee colony algorithm[J]. *Journal of Molecular Modeling*, 2015, 21(10): 261. (JCR Q2)

Invited Talks

2021.12	NeurIPS 2021 (Thirty-fifth Conference on Neural Information Processing Systems), Virtual talk.
2020.09	ECCB 2020 (The 19th European Conference on Computational Biology), Virtual talk.
2020.03	CEGS GGR (Centers of Excellence in Genomic Science Seminar), Stanford University, USA.
2019.11	BIBM 2019 (2019 IEEE International Conference on Bioinformatics & Biomedicine), San Diego, USA
2019.07	ISMB 2019 (The 27th Conference on Intelligent Systems for Molecular Biology), Basel, Switzerland
2019.03	BUAFAI 2019 (The First Beijing Universities Academic Forum of Artificial Intelligence), Beijing, China
2017.01	APBC 2017 (The Fifteenth Asia Pacific Bioinformatics Conference), Shenzhen, China

Honers and Awards

2021.06	Excellent Ph.D. Thesis , Tsinghua University (6% of all Ph.D. theses)
2021.06	Excellent Graduates of Tsinghua University , Tsinghua University (5% of all graduate students)
2021.06	Excellent Graduates of Beijing , Beijing Municipal Commission of Education (5% of all graduate students)
2020.10	Friends of Tsinghua - Mingwei Zhang Scholarship , Tsinghua University
2020.08	ECCB Fellowship , International Society for Computational Biology
2019.10	The First Class Scholarship , Tsinghua University
2019.07	CSC Scholarship , China Scholarship Council
2019.04	ISMB Travel Fellowship , International Society for Computational Biology
2018.10	National Scholarship , Ministry of Education of China (7 out of all Ph.D. students in Department of Automation, Tsinghua University)
2017.10	The First Class Scholarship , Tsinghua University
2016.06	Excellent Graduates of Beijing , Beijing Municipal Commission of Education
2016.06	Excellent Graduates of Beihang University , Beihang University
2016.05	Rui An First Prize Scholarship , Rui An Inc
2015.10	National Encouragement Scholarship , Ministry of Education of China
2015.06	Microsoft Young Fellowship , Microsoft Research Asia (40 undergraduates among top universities in China)
2014.10	National Encouragement Scholarship , Ministry of Education of China
2013.10	National Encouragement Scholarship , Ministry of Education of China
2012.09	Excellent Freshman Prize , Beihang University

Competitions

- 2021.12 **NeurIPS 2021 Competition, Multimodal Data Integration**, rank 1st out of 2611 submissions in two Joint Embeddings
NeurIPS conference
- 2021.05 **RNA Unpaired Probability Prediction Competition**, rank 3/1012, Baidu AI Studio,
Baidu Inc
- 2019.03 **Liver Cancer Image Diagnose Competition**, rank:2/1397, Digital China Innovation Contest,
DCIC 2019
- 2015.01 **Honorable Mention in COMAP's Mathematical Contest in Modeling**, The American
Mathematical Society
- 2014.10 **1st Prize in NCSMC**(The 6th National College Students Mathematical Competition),
Chinese Mathematical Society
- 2014.09 **National 1st Prize in CUMCM**(Contemporary Undergraduate Mathematical Contest in Modeling),
China Society for Industrial and Applied Mathematics
- 2014.09 **1st Prize in Mathematical Competition**, Beihang University
- 2013.12 **1st Prize in Physical Competition**, Beihang University
- 2013.10 **1st Prize in NCSMC**(The 5th National College Students Mathematical Competition),
Chinese Mathematical Society

Technical Strengths

- **Programming Languages**
Python,C,Shell,R,Matlab
- **Deep learning software stacks**
TensorFlow,Keras,PyTorch,PyTorch Lightning,Theano
- **Miscellaneous**
Git,OpenMP,Slurm,Flask,Apache Web Servers

Teaching Experiences

- 2019.02-2019.06 **Teaching Assistant**, Fundamental Industry Training Center, Tsinghua University
Smart Things and Intelligent Systems, Undergraduate Course
- 2018.09-2019.01 **Teaching Assistant**, Department of Automation, Tsinghua University
Introduction to Artificial Intelligence, Undergraduate Course
- 2017.09-2018.01 **Teaching Assistant**, Department of Automation, Tsinghua University
Introduction to Artificial Intelligence, Undergraduate Course
- 2016.08-2016.09 **Teaching Assistant**, Department of Automation, Tsinghua University
Project of Electronic Circuits, Undergraduate Summer Course

Professional Activities

Member of International Society of Computational Biology (ISCB).

Student member of Institute of Electrical and Electronics Engineers (IEEE).

Independent Reviewer for Journals: *Journal of the American Statistical Association*, *Nature Machine Intelligence*, *Bioinformatics*, *BMC Genomics*, *BMC Bioinformatics*, *Complex & Intelligent Systems*.