

# Qiao Liu – Curriculum Vitae

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<b>Address</b>	James H. Clark Center Stanford, CA, 94305	<b>Mobile Phone</b>	(+1)6502835489 (+86)18810226334
<b>Website</b>	<a href="http://liuqiao.me/">http://liuqiao.me/</a>	<b>Email</b>	liuqiao@stanford.edu

## 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

<b>2021.06- Present</b>	Postdoctoral Scholar - Stanford University, Stanford, USA Department of Statistics, advised by Prof. <b>Wing Hung Wong</b> (NAS member) Developing ML algorithms for solving problems in general statistics and computational biology.
<b>2022.01- Present</b>	Part-time Researcher - Zhejiang Laboratory, Zhejiang, China Researcher of AI algorithms in the application of drug development.
<b>2021.06- 2022.02</b>	Part-time Consultant - Shanghai Artificial Intelligence Laboratory, Shanghai, China Research consulting in the combination of AI, computational biology, and medical health care.

## Education

<b>2019.09- 2021.06</b>	Joint Ph.D. student - Stanford University, Stanford, USA Department of Statistics, advised by Prof. <b>Wing Hung Wong</b> (NAS member) Developing machine learning algorithms with applications in both general statistics and biomedical data
<b>2016.09- 2021.06</b>	Ph.D. candidate in <b>Control Science and Engineering</b> - Tsinghua University, Beijing, China Department of Automation, advised by tenured Assoc. Prof. <b>Rui Jiang</b> Tsinghua National Laboratory for Informatics and Technology (TNLIST)
<b>2015.08- 2016.01</b>	Exchange Student - Lund University, Lund, Sweden Department of Computer Science GPA: <b>5.0</b> /5.0
<b>2012.09- 2016.06</b>	Bachelor in <b>Engineering</b> - Beihang University, Beijing, China ShenYuan Honors College (Special Pilot Class, formerly a.k.a. School of Advanced Engineering) GPA: <b>91.5</b> /100, Rank: <b>2</b> /50

## Internships

<b>2019.06- 2019.09</b>	Research Intern - SenseTime Inc, Beijing, China AI+Healthcare, mentored by Dr. <b>Mu Zhou</b> Developing ML algorithms for drug sensitivity prediction and novel drug discovery
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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**<sup>\*</sup>, Rui Jiang<sup>\*</sup> and Wanwen Zeng<sup>\*</sup>. 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<sup>\*</sup>, Dimitri N. Metaxas<sup>\*</sup>. 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<sup>\*</sup>, Rui Jiang<sup>\*</sup>. DeepCAGE: incorporating transcription factors in genome-wide prediction of chromatin accessibility[J]. *Genomics, Proteomics & Bioinformatics*, 2022.
2. **Qiao Liu**, Jiaze Xu, Rui Jiang<sup>\*</sup>, Wing Hung Wong<sup>\*</sup>. 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<sup>\*</sup>, Wing Hung Wong<sup>\*</sup>. 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<sup>\*</sup>, Mu Zhou<sup>\*</sup>. 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<sup>\*</sup>. 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 Biologythe 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>, Xiaolu Fei<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>. 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<sup>\*</sup>, Changjun Zhou<sup>\*</sup>. 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

<b>2022.08</b>	<b>JSM 2022</b> (Joint Statistical Meetings), Wasington DC, USA.
<b>2021.12</b>	<b>NeurIPS 2021</b> (Thirty-fifth Conference on Neural Information Processing Systems), Virtual talk.
<b>2020.09</b>	<b>ECCB 2020</b> (The 19th European Conference on Computational Biology), Virtual talk.
<b>2020.03</b>	<b>CEGS GGR</b> (Centers of Excellence in Genomic Science Seminar), Stanford University, USA.
<b>2019.11</b>	<b>BIBM 2019</b> (2019 IEEE International Conference on Bioinformatics & Biomedicine), San Diego, USA
<b>2019.07</b>	<b>ISMB 2019</b> (The 27th Conference on Intellegent Systems for Molecular Biology), Basel, Switzerland
<b>2019.03</b>	<b>BUAFAI 2019</b> (The First Beijing Universities Academic Forum of Artificial Intelligence), Beijing, China
<b>2017.01</b>	<b>APBC 2017</b> (The Fifteenth Asia Pacific Bioinformatics Conference), Shenzhen, China

## Honers and Awards

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

**2012.09**      **Excellent Freshman Prize**, Beihang University

## Competitions

**2021.12**      **NeurIPS 2021 Competition, Multimodal Data Integration**, rank 1<sup>st</sup>/2611 in two Joint Embedding tasks  
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**      **1<sup>st</sup> Prize in NCSMC**(The 6<sup>th</sup> National College Students Mathematical Competition),  
Chinese Mathematical Society

**2014.09**      **National 1<sup>st</sup> Prize in CUMCM**(Contemporary Undergraduate Mathematical Contest in Modeling),  
China Society for Industrial and Applied Mathematics

**2013.10**      **1<sup>st</sup> Prize in NCSMC**(The 5<sup>th</sup> 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*, *Briefings in Bioinformatics*, *BMC Genomics*, *BMC Bioinformatics*, *Engineering Applications of Artificial Intelligence*, *Complex & Intelligent Systems*, *Computational Statistics*