

# Yu Li

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PhD student  
Computer, Electrical and Mathematical Sci. & Eng.  
King Abdullah University of Science & Technology

## Education

<b>King Abdullah University of Science &amp; Technology (KAUST)</b>	Jeddah, Saudi Arabia
Ph.D. in Computer Science (Advisor: Xin Gao)	2017 - Present
<b>King Abdullah University of Science &amp; Technology (KAUST)</b>	Jeddah, Saudi Arabia
M.S. in Computer Science (GPA: 3.89/4.0)	2015 - 2016
<b>University of Science &amp; Technology of China (USTC)</b>	Hefei, China
B.S. (Honored) in Biosciences (GPA: 3.87/4.3)	2011 - 2015

## Research Interests

- Biomolecular function prediction with deep learning
- Biological sequence analysis and Nanopore sequencing
- Biomolecular structure reconstruction and prediction
- Bio-network analysis with graph embedding
- Machine learning for health care
- Catastrophic forgetting and incremental learning

## Publications

**Journal:**(\*Equal contribution, †Co-corresponding)

24. J Lam\*, **Y Li\***, L Zhu<sup>†</sup>, R Umarov, H Jiang, A Heliou, F Sheong, T Liu, Y Long, Y Li, L Fang, R Altman, W Chen<sup>†</sup>, X Huang<sup>†</sup>, X Gao<sup>†</sup>. "A deep learning framework to predict binding preference of RNA constituents on protein surface". *Nature Communications*. (2019). s41467-019-12920-0.
23. G Jia, **Y Li**, H Zhang, I Chattopadhyay, A Jensen, D Blair, L Davis, P Robinson, T Dahln, S Brunak, M Benson, G Edgren, N Cox, X Gao, A Rzhetsky. "Estimating genetic parameters in the absence of genetic data from country-scale health datasets". *Nature Communications*, *accepted*
22. J Lei, G Sheng, P Cheung, S Wang, **Y Li**, X Gao, Y Zhang, Y Wang, X Huang. "Two symmetric Arginine residues play distinct roles in Thermus thermophilus Argonaute DNA guide strand-mediated DNA target cleavage". *Proceedings of the National Academy of Sciences of the United States of America (PNAS)*. (2019). 10.1073/pnas.1817041116.
21. **Y Li**, T Zhang, S Sun<sup>†</sup>, X Gao<sup>†</sup>. "Accelerating Flash Calculation through Deep Learning Methods". *Journal of Computational Physics (JCP)*. (2019). 10.1016/j.jcp.2019.05.028.
20. **Y Li**, C Huang, L Ding, Z Li, Y Pan, X Gao. "Deep learning in bioinformatics: introduction, application, and perspective in big data era". *Methods*. (2019). 10.1016/j.ymeth.2019.04.008.  
**Cover article of the Methods issue: Deep Learning in Bioinformatics.**

19. Z Zou, S Tian, X Gao, **Y Li**. “mlDEEPre: Multi-functional enzyme function prediction with hierarchical multi-label deep learning”. *Frontiers in Genetics*. (2019) 10.3389/fgene.2018.00714.
18. R Umarov, H Kuwahara, **Y Li**, X Gao<sup>†</sup>, V Solovyev<sup>†</sup>. “Promoter analysis and prediction in the human genome using sequence-based deep learning models”. *Bioinformatics*. (2019). 10.1093/bioinformatics/bty1068.
17. U Hameed, C Liao, A Radhakrishnan, F Huser, S Aljedani, X Zhao, A Momin, F Melo, X Guo, C Brooks, **Y Li**, X Cui, X Gao, J Ladury, L Jaremko, M Jaremko, J Li, S, Arold. “H-NS uses an autoinhibitory conformational switch to achieve environment-controlled gene silencing”. *Nucleic Acids Research (NAR)*. (2018). 10.1093/nar/gky1299.
16. Zhihao Xia, **Y Li**, B Zhang, Z Li, Y Hu, W Chen<sup>†</sup>, X Gao<sup>†</sup>. “DeeReCT-PolyA: a robust and generic deep learning method for PAS identification”. *Bioinformatics*. (2018). 10.1093/bioinformatics/bty991.
15. **Y Li**, R Han, C Bi, M Li, S Wang<sup>†</sup>, X Gao<sup>†</sup>. “DeepSimulator: a deep simulator for nanopore sequencing”. *Bioinformatics*. (2018). 10.1093/bioinformatics/bty223.
14. V Kordopati, A Salhi, R Razali, A Radovanovic, F Tifratene, M Uludag, **Y Li**, A Bokhari, A AlSaieedi, A Raies, C Neste, M Essack, V Bajic. “DES-Mutation: System for Exploring Links of Mutations and Diseases”. *Scientific Reports*. (2018). 10.1038/s41598-018-31439-w.
13. **Y Li**<sup>\*</sup>, F Xu<sup>\*</sup>, F Zhang, P Xu, M Fan, L Li, X Gao<sup>†</sup>, R Han<sup>†</sup>. “DLBI: Deep learning guided Bayesian inference for structure reconstruction of super-resolution fluorescence microscopy”. *Bioinformatics*. (2018). 10.1093/bioinformatics/bty241.
12. S Wang, S Fei, Z Wang, **Y Li**, J Xu, F Zhao, X Gao. “PredMP:a web server for de novo prediction of membrane protein”. *Bioinformatics*. (2018). 10.1093/bioinformatics/bty684.
11. R Han, X Wan, L Li, A Lawrence, P Yang, **Y Li**, S Wang, F Sun, Z Liu, X Gao, F Zhang. “AuTom-dualx: a toolkit for fully automatic alignment of dual-axis tilt series with simultaneous reconstruction”. *Bioinformatics*. (2018). 10.1093/bioinformatics/bty620.
10. R Han, **Y Li**, X Gao, S Wang. “An accurate and rapid continuous wavelet dynamic time warping algorithm for end-to-end mapping in ultra-long nanopore sequencing”. *Bioinformatics*. (2018). 10.1093/bioinformatics/bty555.
9. **Y Li**, S Wang, R Umarov, B Xie, M Fan, L Li, X Gao. “DEEPre: sequence-based enzyme EC number prediction by deep learning”. *Bioinformatics*. (2017). 10.1093/bioinformatics/btx680.
8. H Dai, R Umarov, H Kuwahara, **Y Li**, L Song<sup>†</sup>, X Gao<sup>†</sup>. “Sequence2Vec: a novel embedding approach for modeling transcription factor binding affinity landscape”. *Bioinformatics*. (2017). 10.1093/bioinformatics/btx480.
7. S Wu, D Wang, J Liu, Y Feng, J Weng, **Y Li**, X Gao, J Liu, W Wang. “The dynamic multisite interactions between two intrinsically disordered proteins”. *Angewandte Chemie*. (2017). 10.1002/anie.201701883.
6. X Li, Q Tao, Y Fang, C Cheng, Y Hao, J Qi, **Y Li**, W Zhang, Y Wang, X Zhang. “Reward sensitivity predicts ice cream-related attentional bias assessed by inattention blindness”. *Appetite*. (2015). 10.1016/j.appet.2015.02.010.

**Conference:**(\*Equal contribution, <sup>†</sup>Co-corresponding)

5. L Ding, M Yu, L Liu, F Zhu, Y Liu, **Y Li**, L Shao. Two Generator Game: Learning to Sample via Linear Goodness-of-Fit Test. *Thirty-third Conference on Neural Information Processing Systems (NeurIPS-19)*
4. L Ding, Z Liu, **Y Li**, S Liao, Y Liu, P Yang, G Yu, L Shao, X Gao. "Linear Kernel Tests via Empirical Likelihood for High Dimensional Data". *The Thirty-Third AAAI Conference on Artificial Intelligence (AAAI-19)*
3. L Ding, S Liao, Y Liu, **Y Li**, P Yang, Y Pan, C Huang, L Shao, X Gao. "Approximate Kernel Selection with Strong Approximate Consistency". *The Thirty-Third AAAI Conference on Artificial Intelligence (AAAI-19)*
2. **Y Li**<sup>\*</sup>, F Xu<sup>\*</sup>, F Zhang, P Xu, M Fan, L Li, X Gao<sup>†</sup>, R Han<sup>†</sup>. "DLBI: Deep learning guided Bayesian inference for structure reconstruction of super-resolution fluorescence microscopy". *The Twenty-Sixth Conference on Intelligent Systems for Molecular Biology (ISMB-18)*
1. R Han, **Y Li**, X Gao, S Wang. "An accurate and rapid continuous wavelet dynamic time warping algorithm for end-to-end mapping in ultra-long nanopore sequencing". *The Seventeenth European Conference on Computational Biology (ECCB-18)*

**Under review & Preprint**(\*Equal contribution, <sup>†</sup>Co-corresponding)

6. X Chen<sup>\*</sup>, **Y Li**<sup>\*</sup>, R Umarov, X Gao, L Song, RNA Secondary Structure Prediction By Learning Unrolled Algorithms, *ICLR2020*, *under review*
5. **Y Li**<sup>\*</sup>, S Wang<sup>\*</sup>, X Gao, DeepSimulator1.5: a more powerful, quicker and lighter simulator for Nanopore sequencing, *Bioinformatics*, *under review*
4. H Li<sup>\*</sup>, S Tian<sup>\*</sup>, **Y Li**<sup>\*</sup>, R Tan, Y Pan, C Huang, Y Xu, and X Gao, Modern Deep Learning in Bioinformatics, *Journal of Molecular Cell Biology*, *under review*
3. **Y Li**, Z Li, C Huang, L Ding, Y Pan, X Gao. "SupportNet: solving catastrophic forgetting in class incremental learning with support data". *arxiv.org/abs/1806.02942*
2. **Y Li**, L Ding, X Gao. "On the decision boundary of deep neural network". *arxiv.org/abs/1808.05385*
1. **Y Li**, H Kuwahara, P Yang, L Song<sup>†</sup>, X Gao<sup>†</sup>. "PGCN: Disease gene prioritization by disease and gene embedding through graph convolutional neural networks". *bioRxiv 532226*, 2019

## Experience

<b>Le Song's group, Georgia Tech</b>	Atlanta, USA
Visiting student (Advisor: Le Song)	Fall, 2018
<b>LAMDA, Nanjing University</b>	Nanjing, China
Visiting student (Advisor: Zhi-Hua Zhou)	Summer, 2018
<b>Huang Research Group, HKUST</b>	Hong Kong, China
Visiting student (Advisor: Xuhui Huang)	Summer, 2017
<b>Cognitive Psychology Lab, USTC</b>	Hefei, China
Research assistant (Advisor: Xiaochu Zhang)	2014 - 2015
<b>New Oriental School</b>	Hefei, China
Intern	Winter, 2014

Peking University  
Summer school  
SIBS & IBP, CAS  
Intern

Beijing, China  
Summer, 2014  
Shanghai & Beijing, China  
Summer, 2012

## Patents

6. PGCN: Disease gene prioritization by disease and gene embedding through graph convolutional neural networks (under application)
5. Continuous Wavelet-Based Dynamic Time Warping Method and System (under application)
4. Incremental learning method through deep learning and support data, WO2019193462A1
3. Deep-learning based structure reconstruction method and apparatus, WO2019145767A1
2. Deepsimulator method and system for mimicking nanopore sequencing, WO2019116119A1
1. System, apparatus, and method for sequence-based enzyme ec number prediction by deep learning, WO2019077494A1

## Professional Activities

### Memberships:

- Association for the Advancement of Artificial Intelligence(AAAI) Member
- International Society for Computational Biology (ISCB) Member

### Talks:

6. October 2019, Towards understanding protein functions through deep learning, KAUST Bioengineering Graduate Seminar, Thuwal, Saudi Arabia
5. October 2019, Towards understanding protein functions through deep learning, SUSTech, Shenzhen, China
4. March 2019, Tutorial on Machine Learning and Hand-on Coding Training, CTPL Workshop on Machine Learning, Thuwal, Saudi Arabia
3. July 2018, DLBI: Deep learning guided Bayesian inference for structure reconstruction of super-resolution fluorescence microscopy, The Twenty-Sixth Conference on Intelligent Systems for Molecular Biology (ISMB-18), Chicago, USA
2. June 2018, Deep learning in bioinformatics, Nanjing University, Nanjing, China
1. December 2017, DEEPre: sequence-based enzyme EC number prediction by deep learning, KAUST Research Conference on Big Data Analyses in Evolutionary Biology, Thuwal, Saudi Arabia

### Reviewer:

- Bioinformatics
- IEEE Transactions on Pattern Analysis and Machine Intelligence (TPAMI)

- International Conference on Research in Computational Molecular (RECOMB-20)
- International Conference on Learning Representations (ICLR-20)
- AAAI Conference on Artificial Intelligence (AAAI-18, 19, 20)
- International Joint Conferences on Artificial Intelligence (IJCAI-18)
- Genomics, Proteomics & Bioinformatics (GPB)
- Scientific Reports
- BMC Genomics
- BMC Bioinformatics
- Frontiers in Genetics
- Fuel
- IEEE International Conference on Bioinformatics and Biomedicine (BIBM-17)
- IEEE/ACM Transactions on Computational Biology and Bioinformatics (TCBB)
- Journal of Bioinformatics and Computational Biology (JBCB)
- Journal of Biomedical Informatics (JBI)
- PLOS ONE
- Proteins: Structure, Function, and Bioinformatics

## Supervised Students

### Master students:

- Wenkai Han
- Longxi Zhou

### Visiting students:

- Siyuan Chen (2019, from UESTC)
- Noura AlRasheed (Summer, 2019, KGSP student from UCSD)
- Longxi Zhou (Summer, 2018, from USTC)
- Ammar Alqatari (Summer, 2018, KGSP student from Stanford)
- Zhongxiao Li (Spring, 2018, from SUSTech)
- Zhenzhen Zou (2018, from CAS)

## Teaching

TA for CS320 (Probabilistic Graphical Models) in Spring 2019.

TA for CS229 (Machine Learning) in Spring 2018.

TA for CS220 (Data Analytics) in Fall 2016, Spring 2017, Fall 2017, Fall 2019.

## Awards & Honours

8. Full scholarship for MS/PhD study (acceptance rate: 3%), 2015, KAUST
7. Certificate of Honor Rank (5% out of all graduates in 2015 at USTC), 2015, USTC
6. TianYi FeiYoung Scholarship, 2014, USTC
5. Shizhang Bei's Fellowship, 2013-2015, USTC
4. National Scholarship (2 out of 73 students), 2013, USTC
3. Xingye Scholarship of Responsibility (2 out of 90 students), 2012, USTC
2. Outstanding Student Scholarship, Golden Prize (5% chance of receiving), 2012, USTC
1. Cyrus Tangs Moral Education Scholarship (2 out of 90 students), 2011-2015, USTC