Yichao Li, Ph.D.

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Research Interest

- Genetics and Epigenetics of Hematopoiesis and Blood Diseases
- New Methods for Genome Editing Studies
- Genomics Data Analysis and Visualization
- Machine Learning and Deep Learning

Education

Ohio University

Ph.D., Electrical Engineering and Computer Science
Ohio University

M.S., Mathematics
M.S., Computer Science

Capital Normal University

2012 - 2015

Patent

• B.S., Biological Science

 Hoffmann, R., Frolov, A., Spiller, S., <u>Li, Y</u>. and Welch, L.R., Universität Leipzig and Ohio University. (2017). Method and means for the non-invasive diagnosis of type ii diabetes mellitus. U.S. Patent Application 15/101,885.

Publications

- Lazzarotto, C. R., Malinin, N. L., <u>Li, Y.</u>, Zhang, R., Yang, Y., Lee, G. H., Cowley, E., He, Y., Lan, X., Jividen, K., Katta, V., Kolmakova, N. G., Petersen, C. T., Qi, Q., Strelcov, E., Maragh, S., Krenciute, G., Ma, J., Cheng, Y., & Tsai, S. Q. (2020). CHANGE-seq reveals genetic and epigenetic effects on CRISPR—Cas9 genome-wide activity. *Nature Biotechnology*.
- Qi, Q., Cheng, L., Tang, X., He, Y., <u>Li, Y.</u>, Yee, T., Shrestha, D., Feng, R., Xu, P., Zhou, X., Pruett-Miller, S. M., Hardison, R. C., Weiss, M. J., & Cheng, Y. (2020). Dynamic CTCF binding directly mediates interactions among cis-regulatory elements essential for hematopoiesis. *Blood*.
- Foster, J. M., Grote, A., Mattick, J., Tracey, A., Tsai, Y. C., Chung, M., Cotton, J. A., Clark, T. A., Geber, A., Holroyd, N., Korlach, J., <u>Li, Y.</u>, Libro, S., Lustigman, S., Michalski, M. L., Paulini, M., Rogers, M. B., Teigen, L., Twaddle, A., ... Ghedin, E. (2020). Sex chromosome evolution in parasitic nematodes of humans. *Nature Communications*.
- Grote, A#., Li, Y#., Liu, C., Voronin, D., Geber, A., Lustigman, S., Unnasch, T. R., Welch, L., & Ghedin, E. (2020). Prediction pipeline for discovery of regulatory motifs associated with brugia Malayi molting. PLoS Neglected Tropical Diseases.
- Lee, K. Y., Sharma, R., Gase, G., Ussar, S., <u>Li, Y.</u>, Welch, L., Berryman, D. E., Kispert, A., Bluher, M.,
 & Kahn, C. R. (2017). Tbx15 Defines a Glycolytic Subpopulation and White Adipocyte Heterogeneity. *Diabetes*.
- Spiller, S., Li, Y., Blüher, M., Welch, L., & Hoffmann, R. (2017). Glycated lysine-141 in haptoglobin

- improves the diagnostic accuracy for type 2 diabetes mellitus in combination with glycated hemoglobin HbA1c and fasting plasma glucose. *Clinical Proteomics*.
- Spiller, S., <u>Li, Y.</u>, Blüher, M., Welch, L., & Hoffmann, R. (2018). Diagnostic accuracy of protein glycation sites in long-term controlled patients with type 2 diabetes mellitus and their prognostic potential for early diagnosis. *Pharmaceuticals*.
- <u>Li, Y.,</u> Liu, Y., Juedes, D., Drews, F., Bunescu, R., & Welch, L. (2020). Set cover-based methods for motif selection. *Bioinformatics*.
- <u>Li, Y.,</u> Mullin, M., Zhang, Y., Drews, F., Welch, L. R., & Showalter, A. M. (2020). Identification of Cis-Regulatory Sequences Controlling Pollen-Specific Expression of Hydroxyproline-Rich Glycoprotein Genes in Arabidopsis thaliana. *Plants (Basel, Switzerland)*
- <u>Li, Y.,</u> Shen, X. A., Ewing, R. L., & Li, J. (2018). Terahertz spectroscopic material identification using approximate entropy and deep neural network. *Proceedings of the IEEE National Aerospace Electronics Conference, NAECON*.
- Mason, M. J., Schinke, C., Eng, C. L. P., Towfic, F., Gruber, F., Dervan, A., White, B. S., Pratapa, A., Guan, Y., Chen, H., Cui, Y., Li, B., Yu, T., Chaibub Neto, E., Mavrommatis, K., Ortiz, M., Lyzogubov, V., Bisht, K., Dai, H. Y., ... <u>Multiple Myeloma DREAM Consortium</u>, M. M. D. (2020). Multiple Myeloma DREAM Challenge reveals epigenetic regulator PHF19 as marker of aggressive disease. *Leukemia*
- Hill, S. M., Heiser, L. M., Cokelaer, T., Linger, M., Nesser, N. K., Carlin, D. E., Zhang, Y., Sokolov, A., Paull, E. O., Wong, C. K., Graim, K., Bivol, A., Wang, H., Zhu, F., Afsari, B., Danilova, L. V., Favorov, A. V., Lee, W. S., Taylor, D., ... <u>The HPN-DREAM Consortium</u>, Zi, Z. (2016). Inferring causal molecular networks: Empirical assessment through a community-based effort. *Nature Methods*.

Oral Presentation

Appalachian Region Cell Conference

 Oral presentation: Reverse Engineering of The Human Genome: Predicting Protein-DNA Interactions Across Multiple Cell Types, 2017

Post Presentations

Cold Spring Harbor conference on Systems Biology

• Discovering Gene Regulatory Elements Using Coverage-based Heuristics, 2016

ISCB Great Lakes Bioinformatics Conference

- Homology modeling of extensin peroxidase in Solanum Lycopersicum, 2014
- Epigenetic information improves genome wide motif discovery, 2015
- Motif Discovery in co-regulated DNA sequences, 2016

Statewide Users Group meeting at the Ohio Supercomputer Center

Parallelization of an optimized DNA motif analysis pipeline, with applications in multiple species,
 2018

Professional Services

Ad hoc Reviewer

- IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2015 & 2020
- PLOS Computational Biology, 2016

• The Journal of Open Source Software, 2020

Teaching Experience

Ohio University 2012-2017

Teaching Assistant

- EE 3713, Applied Probability and Statistics for Electrical Engineers.
- CS 2400, Intro to Computer Science I.
- CS 5160, Bioinformatics Tools.
- CS 5170, Data Mining with Applications in the Life Sciences.

Leadership

Big Data Club at Ohio University

2017-2018

President

- Hosted Big Data in Finance Lecture Series, featuring default risk analysis. Promoted blockchain techniques by awarding members with CLH coins.
- Led student teams to learn from Kaggle competitions, including Zillow-2017, Mercari-2018, and YT8m-2018. Also provided data science tutorials and lectures.

Extracurricular Activity

Autonomous Car 2018

• Developed a TensorFlow model on NVDIA TX1, including traffic light, obstacle, grass, and edge detection. Optimized the speed for real-time system. Participated several autonomous car competitions and got a second place (2/10) in drag race at IARRC 2018.

Machine Learning Competition

2018

• Kaggle silver medal, ranked at 18 out of 394, The 2nd YouTube-8M Video Understanding Challenge, 2018.