CURRICULUM VITAE

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Employment

Research Assistant, Department of Agronomy and Horticulture, University of Nebraska-Lincoln 2016-present.

Research Assistant, Genomics and Biotechnology Center, Fujian Agriculture and Forestry University 2013-2016.

Education

Graduate, Agronomy (with James C. Schnable), University of Nebraska-Lincoln, 2016-present.

MS, Bioinformatics (with Haibao Tang), Fujian Agriculture and Forestry University, 2013-2016.

BA, Biotechnology, Henan University of Science and Technology, 2009-2013.

Peer Reviewed Publications

Google Scholar

- 1. **Chenyong Miao**, Jinliang Yang, James C Schnable (2018) Optimizing the identification of causal variants across varying genetic architectures in crops. BIORXIV doi: 10.1101/310391 (Preprint)
- 2. **Chenyong Miao**, Jingping Fang, Delin Li, ... James C Schnable, Haibao Tang (8 total authors) (2018) Genotype-Corrector: improved genotype calls for genetic mapping in F2 and RIL populations. Scientific Reports (Accepted)
- 3. Jisen Zhang, Qing Zhang, Leiting Li, ...**Chenyong Miao**, Ray Ming (11 total authors) (2018) Recent Polyploidization Events in Three *Saccharum* Founding Species. Plant Biotechnology Journal doi: 10.1111/pbi.12962
- 4. Jiandong Bao, Meilian Chen, Zhenhui Zhong, ... Chenyong Miao, ... Baohua Wang, Zonghua Wang (16 total authors) (2017) PacBio Sequencing Reveals Transposable Elements as a Key Contributor to Genomic Plasticity and Virulence Variation in *Magnaporthe oryzae*. Molecular Plant doi: 10.1016/j.molp.2017.08.008
- 5. Jingping Fang, **Chenyong Miao**, Rukai Chen, Ray Ming. (2016) Genome-Wide Comparative Analysis of Microsatellites in Pineapple. Tropical Plant Biology doi: 10.1007/s12042-016-9163-6
- 6. Haibao Tang, Xingtan Zhang, **Chenyong Miao**, Jisen Zhang, Ray Ming, James C Schnable, Patrick S Schnable, Eric Lyons, Jianguo Lu. (2015) ALLMAPS: robust scaffold ordering based on multiple maps. Genome Biology doi: 10.1186/s13059-014-0573-1
- 7. Ray Ming, Robert VanBuren, Ching Man Wai, Haibao Tang, ... Chenyong Miao, ... Robert E Paull, Qingyi Yu (70 total authors). (2015) The pineapple genome and the evolution of CAM photosyntheis. NATURE GENETICS doi: 10.1038/ng.3435

Chenyong Miao

Presentations

o.1 Oral Presentations

1. Chenyong Miao, Jinliang Yang, Schnable JC. (2017) A Bayesian-based GWAS method to identify causal variants in complex polygenic traits. Center for Plant Science Innovation Research Group Meeting. Lincoln, Nebraska.

0.2 Poster Presentations

- 1. **Chenyong Miao**, Jinliang Yang, Schnable JC. (2017) Comparative GWAS in *Sorghum bicolor* and *Setaria italica*. 59th maize meeting. St. Louis, Missouri.
- 2. **Chenyong Miao**, Jinliang Yang, Schnable JC. (2017) Comparative GWAS in *Sorghum bicolor* and *Setaria italica*. UNL Spring Research Fair. Lincoln, Nebraska.
- 3. Chenyong Miao, Piyush Pandey, Zhikai Liang, ... Schnable JC. (2018) Analysis of sorghum timeseries phenotype data using nonparametric curve fitting and machine learning. Phenome 2018. Tucson, Arizona.
- 4. **Chenyong Miao**, Jinliang Yang, Schnable JC. (2018) Large-scale simulation studies enabled by HPC reveal the powers of GWAS approaches in dissecting highly polygenic traits in crop species. Supercomputing and Life Sciences Symposium. Lincoln, Nebraska.
- 5. **Chenyong Miao**, Jinliang Yang, Schnable JC. (2018) Large-scale simulation studies enabled by HPC reveal the powers of GWAS approaches in dissecting highly polygenic traits in crop species. Nebraska Plant Breeding Symposium. Lincoln, Nebraska.