

## Contact Information

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# Dustin Mayfield-Jones

## Education

Current Graduate Student  
Dept. of Plant Biology  
University of Illinois Urbana-Champaign

University of Missouri-Columbia  
Masters in Biology  
Graduation Date: May 2014 (GPA: 4.0)

Truman State University  
Bachelor of Science in Biology *summa cum laude*  
Graduation Date: December 2009 (GPA: 3.91)

## Research Experience

*Summer 2017-Current*

*Dr. Andrew Leakey*

Distinguished Illinois Fellow Dept. of Plant Biology, University of Illinois Urbana-Champaign

- Phenotyping leaf epidermis by optical tomography and computer vision to evaluate stomatal patterning across natural diversity and transgenic lines of sorghum and *Setaria*
- Assisting screening of field experiments to evaluate water-use efficiency traits in sorghum and maize

*Summer 2014-Summer 2017*

*Dr. Thomas Brutnell*

Senior Laboratory Technician Donald Danforth Plant Science Center; Saint Louis, MO

- Co-invented, engineered, and programmed scalable phenotyping robot for controlled watering experiments in greenhouses (Dr. M. Feldman awarded USDA –NIFA I-FAST Challenge Award for further development and commercialization)
- Built next generation RNA-sequencing libraries along the leaf developmental gradient in grasses (maize, *Setaria*, rice, and non-model grasses) and optimized Brutnell Lab RNA-seq protocol
- Calibrated, maintained, and programmed Hamilton robotics for automation of RNA-seq library construction with a technical team
- Managed and organized reagents, samples, and datasets for multiple projects
- Coordinated sequencing efforts among multiple projects
- Executed both formal and informal training sessions
- Performed high-throughput RNA and DNA extractions

*Summer 2010-Spring 2014*

*Dr. J. Chris Pires*

Life Science/NIH T32 Fellow Division of Biological Sciences; University of Missouri-Columbia

**Thesis: “Occurrence and Implications of Biological Network Evolution Following Polyploidy”**

- First-authored two published papers reviewing implications of polyploidy in regard to stages following whole genome duplication, adaptive traits like flowering time, and cancer
- Co-authored paper in which I isolated DNA, made Illumina sequencing libraries, assembled plastid genomes, and tested assembly software
- Collaborated on sequencing projects of various scales, from plastid to nuclear genomes by organizing and inventorying RNA, DNA, and plant/animal tissues, making sequencing libraries, teaching others to construct libraries, and managing and transferring sequencing data
- Developed and tested protocols for next generation sequencing, RNA extraction, and seed storage
- Ordered and maintained laboratory and greenhouse materials and seed from vendors and USDA
- Performed bioinformatic and scientific computing such as plastid genome assembly with NextGENe and CAP3 software, genome annotation with Geneious, programming with Linux command line and

Python, sharing of large sequence data sets via SFTP, and website development and maintenance with WordPress and Google business accounts

- Hired, trained, scheduled and managed student workers and earned certificate in the MU Entering Mentoring program
- Co-organized Interdisciplinary Plant Group Seminar Series by inviting, scheduling, and hosting international investigators while representing the organization
- Co-developed and taught workshop on Next Generation Sequencing at the Botany 2012 conference with Drs. Aaron Liston and Chris Pires' Labs

*Spring 2009-Fall 2009*

*Drs. Brent Buckner & Diane Janick-Buckner*

Student Research Assistant

Department of Biology; Truman State University

- Morphologically characterized two maize developmental mutants using light and scanning electron microscopy, macroscopic photo-documentation, RT-PCR, qPCR, and paraffin sectioning of shoot apical meristems and leaf tissue with Toluidine Blue O and FastGreen staining
- Curated value-added annotation of differentially regulated maize genes using bioinformatics tools including BLASTX, InterProScan, gene prediction with MaizeSequence.org, SDSC Biology Workbench, and literature searches

*Summer 2008-Fall 2008*

*Dr. Karen Cone*

Student Research Assistant

Division of Biological Sciences; University of Missouri-Columbia

“Characterization of a Recombinant Inbred Population for Quantitative Trait Locus mapping”

- Created genetic map with PCR-based markers and MapMaker/QTL software package
- Poured and analyzed >100 agarose gels and collected >20,000 genotype data points

“Effect of *rmr1* on *P11-Blotched*”

- Examined the effect of the epigenetic modifier *rmr1* on *P11-Blotched*, an epigenetically regulated maize gene that activates production of purple anthocyanin pigments

*Spring 2007-Fall 2008*

*Dr. John Ma*

Student Research Assistant

Department of Biology; Truman State University

“The Role of Dark Septate Endophytes in Water Stress Responses of the Forage Grass *Bouteloua gracilis*”

- Optimized growth conditions for *Bouteloua gracilis* and harvested caryopsis for experimentation
- Mentored student lab workers

## Publications

Submitted

- Huang P, Wilson M, **Mayfield-Jones D**, Coneva V, Frank M, Chitwood D. (2017) The evolution of Western tonality: a corpus analysis of 24,000 songs from 190 composers over six centuries. PLOSOne.
- Springer NM, Anderson SN, Andorf CM, Ahern KR, Bai F, Barad O, Barbazuk WB, Bass HW, Baruch K, Ben-Zvi G, Buckler ES, Bukowski R, Campbell MS, Cannon EKS, Chomet P, Dawe RK, Davenport R, Dooner HK, Du LH, Du C, Easterling KA, Gault C, Guan JC, Hunter CT, Jander G, Jiao Y, Koch KE, Kol G, Kollner T, Kudo T, Li Q, Lu F, **Mayfield-Jones D**, Mei W, McCarty DR, Noshay J, Portwood II JL, Ronen G, Settles AM, Shem-Tov D, Shi J, Soifer I, Stein JC, Suzuki M, Vera DL, Vollbrecht E, Vrebalov JT, Ware D, Wei S, Wimalanathan K, Woodhouse MR, Xiong W, Brutnell TP. (2017) The W22 genome: a foundation for maize functional genomics and transposon biology. Nature Genetics.

Published

- Burke SV, Wysocki WP, Zuloaga FO, Craine JM, Pires JC, Edger PP, **Mayfield-Jones D**, Clark LG, Kelchner SA, Duvall MR. (2016) Evolutionary relationships in Panicoid grasses based on plastome phylogenomics (Panicoidae; Poaceae). BMC Plant Biology
- Barrett CF, Baker WJ, Comer JR, Conran JG, Lahmeyer SC, Leebens-Mack JH, Li J, Lim GS, **Mayfield-Jones DR**, Perez LG, Medina J, Pires JC, Santos C, Stevenson DW, Zomlefer WB, Davis JI (2016). Plastid genomes exhibit extensive evolutionary rate variation, and provide support for deep phylogenetic relationships in palms and other commelinid monocots. New Phytologist.

- Cotton JL, Wysocki WP, Clark LG, Kelchner SA, Pires JC, Edger PP, **Mayfield-Jones D**, Duvall MR (2015). Resolving deep relationships of PACMAD grasses; a phylogenomic approach. BMC Plant Biology.
- Saarela JM, Wysocki WP, Barrett C, Soreng RJ, Davis JI, Clark LG, Kelchner SA, Pires JC, Edger PP, **Mayfield DR**, Duvall, MR (submitted). Plastid phylogenomics of the cool-season grass subfamily Pooideae (Poaceae). AoB PLANTS.
- Barrett CF, Freudenstein JV, Li J, **Mayfield-Jones D**, Perez L, Pires JC, Santos C. (2014). "Investigating the path of plastic genome degradation in an early-transitional clade of fungus-eating orchids, and implications for parasitic angiosperms." MBE
- Wysocki WP, Clark LG, Kelchner SA, Burke SV, Pires JC, Edger PP, **Mayfield-Jones D**, Triplett JK, Columbus JT, Ingram AL, Duvall MR (2014). "A Multi-Step Comparison of Short-Read Full Plastome Sequence Assembly Methods in Grasses." Taxon
- Edger PP, Tang M, Bird KA, **Mayfield-Jones D**, Conant GC, Mummenhoff K, Koch MA, Pires JC (2014). "Secondary Structure Analyses of the nuclear rRNA Internal Transcribed Spacers and Assessment of its Phylogenetic Utility across the Brassicaceae (Mustards)." PLOS ONE
- **Mayfield-Jones D**, Washburn JD, Arias T, Edger PP, Pires JC, Conant GC (2013). "Watching the grin fade: Tracing the effects of polyploidy on different evolutionary time scales." Sem Cell & Dev Bio.
- Steele PR, Hertweck KL, **Mayfield D**, McKain MR, Leebens-Mack J, Pires JC (2012). "Quality and quantity of data recovered from massively parallel sequencing: Examples in Asparagales and Poaceae." Am J Bot.
- **Mayfield D**, Chen ZJ, Pires JC (2011). "Epigenetic regulation of flowering time in polyploids." Curr Opin Plant Biol 14(2): 174-178.

### Curriculum Development and Teaching Experience

*Spring 2008-Fall 2009*

*Ms. Maureen McHale*

Instructor of "Ethics of Food" 320

Departments of Biology & Agronomy; Truman State University

- Designed and taught course that surveyed moral principles in relation to food via an interdisciplinary approach
- Areas taught included ecology, economics, agriculture/food technology, theology, and philosophy

### Leadership Experience

*2011- 2014*

University of Missouri-Columbia

Biology Graduate Student Organization

- Vice President • Technology and Website Chair
- Interdisciplinary Plant Group
- Seminar Series Co-Organizer • Technology and Website Chair