

## SCHEDULE OF EVENTS

**Talks will be held in the Grand Ballroom  
Posters will be displayed in Midway West**

### Wednesday, March 5, 2025

8:30 AM – 6:00 PM	<b>OPTIONAL PRE-CONFERENCE WORKSHOPS</b>	
8:30 AM - 11:30 AM	<b>Maize Crop Germplasm Committee</b>	Missouri History Museum
1:00 PM – 6:00 PM	<b>Corn Breeding Research (Day 1)</b>	Midway 1 & 2

### Thursday, March 6, 2025

9:00 AM – 6:00 PM	<b>OPTIONAL PRE-CONFERENCE WORKSHOPS</b>	
8:00 AM – 5:00 PM	<b>Corn Breeding Research (Day 2)</b>	Midway 3 & 4
9:00 AM - 4:00 PM	<b>Development &amp; Cell Biology Workshop</b>	Midway 1 & 2
9:30 AM - 3:00 PM	<b>Bayer Tour</b>	Bus pick-up
12:00 PM - 2:00 PM	<b>39N Lunch and Danforth Center Tour</b>	1001 N. Warson Road St. Louis, MO 63132
2:00 PM - 5:00 PM	<b>Computational Resources Workshop</b>	Conductor Room
3:00 PM – 9:30 PM	<b>REGISTRATION</b>	Depot Registration Office
3:00 PM – 6:00 PM	<b>POSTER HANGING</b>	Midway West
5:00 PM – 5:45 PM	<b>Travel awardees / mentors meet &amp; greet</b>	Jeffersonian / Knickerbocker
6:00 PM – 7:00 PM	<b>DINNER</b>	Midway West & Pegram

## **Thursday, March 6, 2025 (continued)**

7:00 PM – 9:00 PM	<b>SESSION 1 – WELCOME / KEYNOTE/ GENE REGULATION</b> Chair: Sherry Flint-Garcia	
7:00 PM	<b>WELCOME AND ANNOUNCEMENTS</b>	Grand Ballroom
7:20 PM	<b>Jonathan Wendel, Iowa State University</b> <i>Genes, jeans, genomes, and the wonders of polyploidy in plants</i>	[KS1]
8:00 PM - 8:20 PM	<b>POSTER LIGHTNING TALKS</b>	
	<b>Yue Liu, Iowa State University</b> <i>Candidate Genes Underlying a Major QTL qshgd1 Causing Spontaneous Haploid Genome Doubling in Maize A427</i>	[L1]
	<b>Vladimir Torres-Rodriguez, University of Nebraska-Lincoln</b> <i>Multi-species transcriptome-wide association studies identify additional genes controlling flowering</i>	[L2]
	<b>Andrea Sama, University of California, San Diego</b> <i>Chemical Imaging Reveals Metabolic Responses to Salt-Stress in Maize Roots</i>	[L3]
	<b>Xiaosa Xu, University of California, Davis</b> <i>A high-resolution, meristem stage-specific single-cell gene expression atlas resolving developmental dynamics in maize inflorescence architecture</i>	[L4]
	<b>Sebastian Mueller, Oregon State University</b> <i>Predictive Modeling of Pollen Fitness Phenotypes from Genome Scale Data Identifies Expression Specificity As a Critically Informative Parameter</i>	[L5]
	<b>Huan Chen, Michigan State University</b> <i>Archaeological Bolivian maize genomes suggest Inca cultural expansion augmented maize diversity in South America</i>	[L6]
	<b>Lukas Würstl, Technical University Munich</b> <i>Natural alleles of the gene lhcb6 shape photosynthesis and key agronomic traits in maize (Zea mays L.) landraces</i>	[L7]
	<b>Hannah Pil, North Carolina State University</b> <i>BZea: A diverse teosinte introgression population for improving modern maize sustainability</i>	[L8]

**Thursday, March 6, 2025 (continued)**

8:20 PM	<b>Maike Stam, University of Amsterdam</b> <i>Vgt1 as enhancer of ZmRap2.7 impacts flowering time and gene regulatory networks involved in jasmonate signaling in maize</i>	[T1]
8:40 PM	<b>Ankush Sangra, University of Georgia</b> <i>Decoding a complex distal non-coding QTL at TEOSINTE BRANCHED 1</i>	[T2]
9:00 PM – 12:00 AM	<b>INFORMAL POSTER VIEWING &amp; HOSPITALITY</b>	Midway West
9:00 PM – 10:30 PM	<b>INFORMAL CORN HOLE PLAY</b>	Prefunction Space Grand Ballroom

## **Friday, March 7, 2025**

7:00 AM – 8:00 AM      **BREAKFAST**      Midway West

7:30 AM – 12:30 PM      **REGISTRATION**      Depot Registration  
Office

8:15 AM – 10:15 AM      **SESSION 2 – MODELING CORN**  
Chair: Hank Bass

8:15 AM      **ANNOUNCEMENTS**      Grand Ballroom

8:30 AM      **Jingjing Zhai, Cornell University**      [T3]  
*Cross-species modeling of plant genomes at single  
nucleotide resolution using a pre-trained DNA  
language model*

8:50 AM      **Diana Ruggiero, Oregon State University**      [T4]  
*Quantitative genetics of leaf vascular density in maize*

9:10 AM      **Erin Farmer, Cornell University**      [T5]  
*Integrating proximal sensing modalities for enhanced  
prediction of agronomically important crop traits*

9:30 AM      **Lucas Batista & Jacob Washburn, Kansas State  
University & USDA-ARS**      [T6]  
*Crowdsourcing phenotype prediction: Results from  
the 2024 G2F prediction competition.*

9:50 AM - 10:15 AM      **POSTER LIGHTNING TALKS**

**Jacob Kelly, University of Missouri**      [L9]  
*Speed Breeding Fast-Flowering Mini-Maize*

**Thanduanlung Kamei, University of Delaware**      [L10]  
*SBP mutants have an expanded competence zone for brace  
root initiation*

**Katy Guthrie, University of Minnesota**      [L11]  
*Teaching Scientific Writing Alongside the Scientific Method  
in an Introductory Plant Biology Lab*

**Joseph DeTemple, Iowa State University**      [L12]  
*Gene expression and circadian rhythm differences between  
temperate and tropical maize inbreds in response to  
photoperiod*

## **Friday, March 7, 2025 (continued)**

	<b>Manisha Munasinghe, University of Minnesota</b> <i>Structural Variation has a Limited Role in Influencing Genome-Wide Differential Gene Expression Patterns in Maize</i>	[L13]
	<b>Mohamed El-Walid, Cornell University</b> <i>Genomic Assembly and Analysis of Fast-Flowering Mini-Maize</i>	[L14]
	<b>Christopher Benson, Ohio State University</b> <i>Resolving Maize Domestication and Subpopulation Divergence Using Long Terminal Repeat Retrotransposons</i>	[L15]
	<b>Xuelian Du, University of Bonn</b> <i>BonnMu – A resource for functional genomics in maize (Zea mays L.)</i>	[L16]
10:15 AM - 10:45 AM	<b>BREAK</b>	Prefunction Space Grand Ballroom
10:45 AM – 12:15 PM	<b>SESSION 3 – EDUCATION, COMMUNITY, AND OUTREACH</b> Chair: Brandi Sigmon	
10:45 AM	<b>ACKNOWLEDGE TRAVEL AWARDEES</b>	Grand Ballroom
10:55 AM - 11:10 AM	<b>POSTER LIGHTNING TALKS</b>	
	<b>Jason Lynn, Cold Spring Harbor Laboratory</b> <i>AGO2 and AGO3 regulate RNAi fidelity by suppressing RNA-directed DNA methylation</i>	[L17]
	<b>Dafang Wang, Hofstra University</b> <i>Mechanisms of Small RNA-Induced Epigenetic Silencing of Ac Transposons in Maize</i>	[L18]
	<b>Vinay Chaudhari, Donald Danforth Plant Science Center</b> <i>Predicting end-of-season Sorghum biomass from seedling-stage traits</i>	[L19]
	<b>Olivia Haley, USDA-ARS, ORISE</b> <i>Comparing the performance of protein folding models AlphaFold, ESMFold, and Boltz for classical genes in maize</i>	[L20]
	<b>Zong-Yan Liu, Cornell University</b> <i>ReelGene2: A Large Language Model for Single Base Pair Precision Gene Annotation in Diverse Plant Genomes</i>	[L21]

## **Friday, March 7, 2025 (continued)**

11:10 AM	<b>Vivian Bernau, USDA Plant Introduction</b> <i>Ensuring the future of maize: A call for collaborative action</i>	[T7]
11:30 AM	<b>Helen Anne Curry, Georgia Tech</b> <i>Input, Insurance, Objective: Reflections on diversity from the history of crop science</i>	[KS2]
12:15 PM - 1:15 PM	<b>LUNCH</b> Special table at lunch w/ Helen Anne Curry	Midway & Pegram
	Travel Awardee/Mentor Networking Lunch	Midway 1 & 2
	MGC BoD and MGAC Lunch	Midway 3 & 4
1:30 PM - 4:30 PM	<b>POSTER SESSION 1</b>	Midway West
1:30 PM - 3:00 PM	<i>Presenters should be at odd-numbered posters</i>	
3:00 PM - 4:30 PM	<i>Presenters should be at even-numbered posters</i>	

Beverages will be available from 2:30 to 4:00 PM in Midway West

### **4:40 PM – 6:00 PM    SESSION 4 – MAIZE UNDER STRESS** Chair: Melissa Draves

4:40 PM	<b>Veronica Perez, Cornell University</b> <i>Translational and proteomic analysis of cold-stressed maize reveals ribosomal protein families involved in cold response and tolerance</i>	[T8]
5:00 PM	<b>Fausto Rodríguez-Zapata, North Carolina State University</b> <i>Introgression of a Mexican highland chromosomal inversion into temperate maize accelerates flowering, promotes growth, and modulates a cell proliferation gene network.</i>	[T9]
5:20 PM	<b>Marie-Laure Martin, INRAE</b> <i>Integration of phenomic, proteomic, and genomic data into a multi-scale network unravels missing heritability for maize response to water deficit</i>	[T10]
5:40 PM	<b>Maggie Woodhouse, USDA-ARS</b> <i>Transcriptional regulation of stress adaptation in maize: identification and functional annotation</i>	[T11]

## **Friday, March 7, 2025 (continued)**

6:00 PM - 7:00 PM	<b>DINNER</b> Bayer Student/Postdoc Dinner	Midway & Pegram Midway 1 & 2
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7:00 PM - 9:00 PM	<b>SESSION 5 - AWARDS</b> Chair: Andrea Eveland
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7:00 PM	<b>Andrea Eveland</b> <i>Introduction to Awards</i>	Grand Ballroom
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7:10 PM	<b>Natalia de Leon</b> <i>Presenting: Cooperator and Leadership Awards</i>
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7:30 PM	<b>Andrea Eveland</b> <i>Presenting: M. Rhoades Early-Career, L. Stadler Mid-Career</i>
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7:50 PM	<b>Andrea Eveland and Wojtek Pawlowski</b> <i>Presenting: R. Emerson Lifetime Awards</i>
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8:10 PM	<b>Marna Yandeau-Nelson</b> <i>McClintock Prize Presentation Introduction</i>
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8:20 PM	<b>Edward Buckler IV, USDA Agricultural Research Service</b> <i>Why do we do maize genetics?</i>	[M1]
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9:00 PM - 12:00 AM	<b>INFORMAL POSTER VIEWING &amp; HOSPITALITY</b>	Midway West
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9:30 PM - 10:00 PM	<b>OPERA BELL BAND</b>	Pegram
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9:00 PM - 10:30 PM	<b>EARLY BRACKETS OF THE CORN HOLE TOURNAMENT</b>	Prefunction Space Grand Ballroom
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## **Saturday, March 8, 2025**

7:00 AM – 8:00 AM      **BREAKFAST**      Midway West & Pegram

8:00 AM – 12:00 PM      **REGISTRATION**      Depot Registration Office

8:15 AM – 10:00 AM      **SESSION 6 – ROOTS & NUTRIENT UPTAKE / KEYNOTE**  
Chair: Rubén Rellán Álvarez

8:15 AM      **ANNOUNCEMENTS**      Grand Ballroom

8:20 AM      **Alexander Liu, Washington University in Saint Louis, Donald Danforth Plant Science Center**      [T12]  
*A Rootless1 knockdown allele affects maize nodal root development, increasing rooting depth, nitrogen uptake efficiency, and grain production in the field*

8:40 AM      **Sylvia Morais de Sousa Tinoco, Embrapa**      [T13]  
*Overexpression of PSTOL1-like genes increases maize root surface area and biomass under low and high phosphorus conditions*

9:00 AM      **Ivan Baxter, Donald Danforth Plant Science Center**      [KS3]  
*You need a real maize geneticist*

9:40 AM – 10:00 AM      **POSTER LIGHTNING TALKS**

**Forrest Li, University of California, Davis**      [L23]  
*Sequencing a seed bank: Assessing the utility of environmental data from CIMMYT traditional varieties for climate-adaptive maize breeding*

**Aimee Schulz, University of Minnesota**      [L24]  
*The molecular evolution of perenniality across the grasses*

**Matthew Wendt, Iowa State University**      [L25]  
*Environmental and Genetic Factors Underlying Maize Cuticular Wax Accumulation Under Drought Stress*

**Wen-Yu Liu, North Carolina State University**      [L26]  
*ZmCER9-Mediated Regulation of Autoactive NLR Proteins and Effector-Triggered Immunity via ERAD Pathway*



## **Saturday, March 8, 2025 (continued)**

	<b>Huda Ansaf, University of Missouri-Columbia</b> <i>Understanding the Role of TOR Signaling and Translational Machinery in Regulating Protein-bound Amino Acid Homeostasis in Maize Kernels</i>	[L27]
	<b>Gwonjin Lee, West Virginia State University</b> <i>Sex-specific patterns of meiotic recombination are determined by maize lines from different climate zones.</i>	[L28]
	<b>Michelle Stitzer, Cornell University</b> <i>Comparative grass genomics reveals explosive genome evolution in maize and its wild relatives</i>	[L29]
	<b>Mohammad Mahmood Hasan, University of Florida</b> <i>mop1 reshapes recombination landscapes by altering DNA methylation and chromatin states at MITEs</i>	[L30]
10:00 AM – 10:30 AM	<b>BREAK</b>	Prefunction Space Grand Ballroom
10:30 AM – 12:30 PM	<b>SESSION 7 – KEYNOTE / CELL DIVISION &amp; MERISTEMS</b> Chair: Sarah Jensen	
10:30 AM	<b>Sióbhán Brady, Howard Hughes Medical Institute, University of California Davis</b> <i>Environmental integration with root cell type development</i>	[KS4]
11:10 AM	<b>Stephanie Martinez, University of California, Riverside</b> <i>Delayed divisions and cell elongation defects influence plant growth in katanin mutants</i>	[T14]
11:30 AM	<b>Fang Xu, Shandong University</b> <i>The EPF-ERECTA ligand-receptor pairs regulate maize shoot and inflorescence architecture in coordination with CLAVATA pathway in maize.</i>	[T15]
11:50 AM	<b>Thu Tran, Cold Spring Harbor Laboratory</b> <i>Catalytic and non-catalytic TREHALOSE-6-PHOSPHATE SYNTHASES (TPSs) interact with RAMOSA3 to control maize development</i>	[T16]

## **Saturday, March 8, 2025 (continued)**

12:10 PM	<b>Alejandro Aragon Raygoza, Iowa State University</b> <i>Exploring the effects of ethylene-related transcription factors during maize shoot development</i>	[T17]
12:30 PM - 1:30 PM	<b>LUNCH</b> <i>Travel awardee lunch with keynote speakers</i> <i>Maize genetics mentoring &amp; networking lunch</i> <i>MGMSC lunch</i>	Midway & Pegram Midway 1 & 2 Midway 3 Midway 4
1:30 PM - 4:30 PM	<b>POSTER SESSION 2</b>	Midway West
1:30 PM - 3:00 PM	<i>Presenters should be at even-numbered posters</i>	
3:00 PM - 4:30 PM	<i>Presenters should be at odd-numbered posters</i>	
Beverages will be available from 2:30 to 4:00 PM in Midway West		
4:30 PM - 6:00 PM	<b>COMMUNITY SESSION</b> <b>Maize Genetics Cooperative</b> <i>Wojtek Pawlowski, MGC BoD Chair</i>	Grand Ballroom
6:00 PM - 7:00 PM	<b>DINNER</b> <i>Corteva Student/Postdoc Dinner</i>	Midway West & Pegram Midway 1 & 2
7:00 PM – 8:20 PM	<b>SESSION 8 – REPRODUCTION / KEYNOTE</b> Chair: Cinta Romay	
7:00 PM	<b>ANNOUNCEMENTS</b>	Grand Ballroom
7:05 PM	<b>Xixi Zheng, University of Regensburg</b> <i>Understanding the Molecular Mechanism of Parthenogenesis in Cereals</i>	[T18]
7:25 PM	<b>Rachel Egger, Syngenta</b> <i>Heat treatment and UBA2 fusions enhance LbCas12a genome editing activity during haploid induction</i>	[T19]
7:45 PM	<b>Elli Cryan, University of California Davis</b> <i>Molecular evolution of the Ga reproductive barriers in maize and related species</i>	[T20]

**Saturday, March 8, 2025 (continued)**

8:05 PM	<b>Doreen Ware, USDA Agricultural Research Service</b> <i>Plant genomes: Understanding their past and managing their future</i>	[KS5]
9:00 PM - 12:00 AM	<b>INFORMAL POSTER VIEWING &amp; HOSPITALITY</b>	Midway West
10:00 PM - 12:00 AM	<b>GAME NIGHT / CORN HOLE TOURNAMENT FINALS</b>	Midway West

## **Sunday, March 9, 2025**

7:00 AM - 8:20 AM     **BREAKFAST**

Midway & Pegram

8:25 AM – 10:20 AM     **SESSION 9 – EPIGENETICS**  
Chair: Katie Murphy

8:25 AM     **ANNOUNCEMENTS**

8:30 AM     **Qi Li, University of Tuebingen, Germany**     [T21]  
*Long-distance retrotransposons direct variable gene imprinting in maize*

8:50 AM     **Xi Cheng, University of Florida**     [T22]  
*Deciphering epigenetic and genetic alterations in a DNA methylation mutant through successive generations of self-fertilization in maize*

9:10 AM     **Hafiza Sara Akram, Florida State University**     [T23]  
*Replication timing uncovers a novel two-compartment arrangement of maize interphase euchromatin*

9:30 AM     **Akwasi Yeboah, University of Florida**     [T24]  
*Determination of Genetic and Epigenetic Regulations of Meiotic Recombination during Domestication in Maize*

9:50 AM - 10:20 AM     **BREAK**

## **Sunday, March 9, 2025 (continued)**

### **10:20 AM – 12:00 PM SESSION 10 – BUILDING A STRONGER MAIZE PLANT**

Chair: Frank Hochholdinger

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| 10:20 AM | <b>Bharath Kunduru, Clemson University</b><br><i>Deciphering genetic architecture of stalk lodging resistance using high-density phenotype map in maize</i>    | [T25] |
| 10:40 AM | <b>Laura Tibbs-Cortes, USDA-ARS</b><br><i>Plasticity and fitness trade-offs in switchgrass revealed by open science and citizen science data</i>               | [T26] |
| 11:00 AM | <b>Qin Yang, Northwest A&amp;F University</b><br><i>Inactivation of a lysine-histidine transporter-1 gene confers southern leaf blight resistance in maize</i> | [T27] |
| 11:20 AM | <b>Marion Pitz, University of Bonn</b><br><i>Regulation of heterosis-associated gene expression complementation in maize hybrids</i>                           | [T28] |
| 11:40 AM | <b>CLOSING REMARKS</b>   |       |
| 12:00 PM | <b>ADJOURNMENT</b>   |       |

### **1:00 PM – 3:00 PM OPTIONAL POST-CONFERENCE WORKSHOP**

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| 1:00 PM - 3:00 PM | <b>Missouri Botanical Garden</b> | Missouri Botanical Garden<br>4344 Shaw Blvd<br>St. Louis, MO 63110 |
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