3rd Multi-Omic Integration for AI Genomic Prediction Breeding Under Different Approaches: Past, Present and Future

PROGRAM

July 07-11, 2025 | Gainesville, FL

https://agronomy.ifas.ufl.edu/moiai/

UF IFAS
UNIVERSITY of FLORIDA

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WI-FI INFORMATION
Network: ufguest

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Name Badge

Your name badge serves as your admission to all networking functions, so be sure to wear it throughout the short course.

UF IFAS
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WELCOME ATTENDEES!

Thank you for joining us for the 3rd Multi-Omic Integration for AI Genomic Prediction Breeding Under Different Approaches: Past, Present and Future Short Course!

Dear Participants,

It is my great pleasure to welcome you to the 3rd Multi-Omic Integration for AI Genomic Prediction Breeding Short Course. We are truly excited to have you join us for this unique opportunity to explore the latest advances at the intersection of genomics, multi-omics, and artificial intelligence in breeding.

This year, we are especially grateful to Dr. David Norton, Vice President of Research at the University of Florida, for his generous support in allowing us to use the NVIDIA-Malachowsky Auditorium. This state-of-the-art facility offers an inspiring environment for learning and collaboration, and we are honored to host this event in such a cutting-edge space.

We are also delighted to announce that, for the first time, our short course is supported by two outstanding sponsors: **Agronomix Software, a CULTURA company**, and **Fall Creek Blueberry Nursery**. Their commitment to advancing research and innovation in agricultural genomics is invaluable, and we deeply appreciate their partnership.

This year's program features an exceptional lineup of speakers, each bringing expertise and insight from across the field. Their contributions will provide you with a comprehensive overview of the latest developments and practical applications in multi-omic integration and AI-driven genomic prediction. We encourage you to engage with our speakers and fellow participants to maximize your learning experience.

Thank you for joining us. We look forward to a stimulating and productive short course!

Sincerely,

Diego Jarquín

On behalf of the organizing committee

GUEST SPEAKERS

Dr. John Davis

Associate Dean for Research
UF/IFAS Office Of The Dean For Research

John leads the research enterprise at UF/IFAS, supporting faculty in Gainesville and at our research and education centers who address challenges facing agriculture, natural resources and human systems. He also oversees operations at the Plant Science Research and Education Unit and the Ordway-Swisher Biological Station.
John Davis specialized in forest ecosystem health, tree-pathogen coevolution, forest genomics, and is tenured in the School of Forest, Fisheries, and Geomatics Sciences. He earned his Ph.D. in plant breeding and genetics-forestry from Michigan State University in 1989 and has been a faculty member in UF/IFAS since 1992. Prior to serving in the Dean for Research Office, he maintained a 60 percent research, 20 percent teaching, and 20 percent administrative appointment since 2011. In March 2016, he transferred to the Dean for Research Office as Associate Dean and in February 2024 he became Interim Dean and Director of Florida Agricultural Experiment Station.

Dr. Jason Ferrell

Professor & Department Chair
UF/IFAS Agronomy Department

Dr. Ferrell joined the University of Florida as an Assistant Professor in 2004 with responsibilities in weed management and herbicide stewardship. In the time since, he has attained the rank of Associate Professor (2009) and Professor (2014). Dr. Ferrell became the Director of the UF'/IFAS Center for Aquatic and Invasive Plants in 2017 where he helps coordinate and promote invasive plant research and extension activities throughout the state. In 2019 to 2024, he served as Director of the UF/IFAS Pesticide Information Office which directs pesticide licensing/testing and extension outreach activities for pesticide safety and compliance. In 2024, he was named Chair of the Agronomy Department and provides direction to 36 faculty programs located across 8 statewide units.

COURSE INSTRUCTORS

Dr. Diego Jarquín

Assistant Professor
UF/IFAS Agronomy
Integration and Application of AI and Omics in Plant Breeding

One of the biggest challenge of the humanity is to ensure the current and future food supply chain of a growing population in a world that is facing more often and extreme environmental variations. The contributions of Dr. Jarquín research program are relevant to the society because these are helping to the development of improved (more productive, resistant to biotic and abiotic stress, etc.) genotypes by the integrating artificial intelligence (AI) methods and multi-omics analyses in plant breeding. Dr. Jarquíjan's program is focused on the development of interpretable AI methods and related disciplines (biostatistics, quantitative genetics, and modeling) that can be applied to large multi-omics data sets for providing solutions to complex plant breeding and plant systems biology questions. More specifically, these developments are oriented to find new ways of driving genetic improvement and biological insights designing and optimizing methods for plant breeding, leveraging information from multiple facets of plant biology-physiology, agronomy, and biochemistry to quantitative genetics and multi-omics (genomics, transcriptomics, proteomics, metabolomics and high throughput phenotyping), and provide novel solutions to unravel the biological basis of complex traits for plant breeding programs.

Dr. Charlie Messina

Professor & Director of Crop Transformation Center
UF/IFAS Horticultural Sciences
Predictive Breeding

Charlie Messina is Professor in the Department of Horticultural Sciences at the University of Florida. His teaching and research program focuses on how to harmonize crop improvement and AI efforts for agricultural systems to regenerate the environment while providing nutrition security, improve human health and adapt to climate change. His team evaluates and applies novel prediction methods and systems within operational breeding programs. During his tenure at Corteva Agriscience he contributed to the development of drought tolerant maize in the U.S. and Brazil, which is now grown in >10MM acres.

Dr. Esteban Ríos

Associate Professor
UF/IFAS Agronomy
Forage Breeding and Genetics

Dr. Ríos research program meets a critical need for society as the demand for animal-based products is exponentially growing, particularly in subtropical and tropical regions where population growth is more pronounced. The U.S. is the world's largest beef producer and cattle production accounts for more than $78 billion in sales annually. However, feed represents the highest cost component of beef production and accounts for 40-60% of production costs. The interest in utilizing improved forages in pastures, versus grain-fed approaches, is driven by economic factors, followed by environmental and social consciousness. Improved forages contribute to reducing greenhouse gas emissions from the livestock sector, in addition to sequestering carbon in soil to mitigate climate change. His specialization in plant breeding and genetics, quantitative genetics, and forage production are the foundation of his research questions to improve yield, nutritive value, abiotic and biotic stress tolerance in forage species.

Dr. Paulino Pérez-Rodríguez

Professor
Statistics and Data Science
Colegio de Postgraduados

Dr. Pérez-Rodríguez did his MS and Doctoral degrees in Statistics from the Colegio de Postgraduados, México. In 2011/12 he completed a 1-year postdoc at the University of Wisconsin-Madison under the supervision of Dr. Daniel Gianola and in 2012/2013 he became a visiting scholar at the Section on Statistical Genetics of the Biostatistics Department of University of Alabama at Birmingham where he worked in collaboration with Dr. Gustavo de los Campos. In 2019/2020 he was a visiting scholar at the Epidemiology and Biostatistics of Michigan State University where he worked in collaboration with Dr. Gustavo de los Campos. Dr. Pérez-Rodríguez has a vast experience as an industry consultant both in plant and animal breeding. Currently he is a Professor of Colegio de Postgraduados in México in the department of Statistics and Data Science. His research interests include Statistical computing, Bayesian statistics, high throughput computing and software development for prediction of complex traits in plants and animals.

Dr. Marcio Resende

Associate Professor
UF/IFAS Horticultural Sciences
Sweet Corn Breeding and Genomics

Dr. Resende has two active breeding programs working on the development of sweet corn inbreds and potato varieties that are aimed at addressing the concerns of the Florida growers. He is also interested in statistical genomics and quantitative genetic research that can increase genetic gains, increase efficiency and/or accelerate the breeding program. His lab has a strong research and training component in different 'omics' with an applied goal. Current projects include the use of metabolomics to predict food sensory panels, the development of genomic selection models to improve breeding and the analysis of transcriptome information for the detection of eQTLs.

Dr. Hsiao-Yi Hung

Quantitative Geneticist
Driscoll's Inc

Dr. Hsiao-Yi Hung is a quantitative geneticist at Driscoll's. He received his Ph.D. in plant breeding and genetics from North Carolina State University. His research focuses on experimental design, quantitative/statistical genetics, genetic evaluation and statistical modeling. He has been working on implementing prediction framework in various crops for more than ten years

Dr. Enid Pérez-Lara

Industry Engagement Manager
Agronomix Software

Dr. Pérez-Lara is a seasoned plant breeder and a passionate advocate for diversity in agricultural sciences. With over 24 years of experience in plant breeding, she has worked across various crops, supporting research, innovation, and technology adoption on a global scale. Currently serving as the Industry Engagement Manager at Agronomix Software, she collaborates with breeding programs worldwide to integrate cutting-edge digital tools that enhance decision-making and efficiency. Beyond her technical expertise, Dr. Perez-Lara is a dedicated champion for women in plant breeding, actively promoting career opportunities for young scientists and fostering international knowledge exchange to drive innovation in the field.

Julián García-Abadillo

Ph.D. Candidate
Quantitative Gen. - AI - G×E

Mr. García-Abadillo Velasco is a biotechnology graduate interested in quantitative genetics, biometrics, and statistics. His academic formation is on Computer Science (MSc. Artificial Intelligence) and he is currrently focusing on improving his programming skills and knowledge on Information Theory, Algorithms and Data Structures to develop applications on real-world problems while also concerned with theoretical issues on Artificial Intelligence such as Knowledge-representation field.

Dr. Ying Zhang

AI and RSE Services Manager
Research Computing
UF Information Technology

Dr. Zhang is the IT Manager for AI and Research Software Engineering Services at Research Computing, a department in Information Technology at University of Florida. She and her team work with faculty research groups and provide consultations in AI research and software development in all research disciplines. The team also design and deliver AI training courses at different competency levels for UF and its partner institutions. She also assists researchers in running high performance computing (HPC) applications with high efficiency on HiPerGator - UF's supercomputing clusters. She has a MS degrees in Computer Science and Remote Sensing.

Dr. Qian Zhao

AI/HPC Support Engineer
Certified University Ambassador @NVIDIA
University of Florida

As an AI Support Engineer at UFIT Research Computing, Chin(Qian) specializes in optimizing AI and data science workloads on high-performance computing (HPC) systems. With expertise in GPU acceleration, parallel computing, and deep learning frameworks, She helps researchers and engineers deploy and fine-tune large-scale AI models for cutting-edge applications. Passionate about performance tuning, scalability, and advancing AI in supercomputing, she thrives on collaborating with teams to push the boundaries of innovation.

Dr. Jorge Hidalgo

Assistant Professor
University of Georgia
Animal Breeding and Genetics

Dr. Jorge Hidalgo focuses on the application and development of statistical methods for large-scale genomic evaluations in livestock populations. The most advanced breeding programs gather millions of pedigrees, genotypes, and phenotypes, leading to more accurate prediction of the animals' genetic merit and a deeper understanding of the genetic basis of production, health, and efficiency traits. The data dimensionality also brings computational challenges, making the development of more efficient statistical methods and their application crucial. Specific research topics include 1) genomic evaluation for crossbred and beef on dairy populations, 2) genetic analysis of categorical traits (diseases, mortality, calving ease, etc.), 3) genetic analysis including social interaction models, and 4) genomic evaluations for heat tolerance.

Vitor Sagae

Ph.D. Candidate
Research Scholar
UF/IFAS Agronomy Department

Mr. Sagae has a Master's degree in Genetics and Breeding from the Federal University of Viçosa, and a BS. in Agronomy from UNICENTRO. He is experienced in plant breeding and seed production, working with maize and soybean in academia and industry. Interested in breeding methods, experimental designs, quantitative genetics, and statistical methods applied to plant breeding.

Dr. Caio Canella Vieira

Assistant Professor
University of Arkansas
Soybean Breeding and Genetics

Dr. Caio Canella Vieira is an Assistant Professor of soybean breeding at the University of Arkansas, currently leading the University's soybean breeding program. Before joining the University of Arkansas, Dr. Canella Vieira obtained his master's and Ph.D. in Plant Breeding, Genetics, and Genomics at the University of Missouri. He was involved in the development and release of over 20 soybean cultivars in Missouri and led the modernization of the soybean breeding program by prioritizing the use of data analytics and prediction models. His program in Arkansas targets the development and release of high-yielding conventional and herbicide-tolerant soybean cultivars with broad biotic and abiotic stressors tolerance, as well as improved seed composition. The program puts efforts into identifying and incorporating economic-important traits from genetically diverse accessions into modern, high-yielding genetic backgrounds. Substantial efforts are also made on maximizing the efficiency of the soybean breeding program through the implementation of genomic prediction and high-throughput phenotyping strategies throughout all stages of the pipeline.

Dr. Huiwen Ju

Solutions Architect
Higher Education & Research
NVIDIA Corporation

Dr. Huiwen Ju is a Solution Architect on the Higher Education & Research team at NVIDIA, helping researchers leverage NVIDIA’s full-stack solution to accelerate their research workloads. Before joining NVIDIA, she earned her PhD in Computational Neuroscience from Georgia State University.

Dr. Julio Isidro y Sánchez

Biotechnology and Plant Biology Department
Centre for Plant Biotechnology and Genomics
University Polytechnic of Madrid

Dr. Isidro y Sánchez is a member of the Computational Systems Biology and Genomics (CsBGP) at the Center of Biotechnology and Genomics in Spain, Madrid. He is leading the research group Crop Genomics and Breeding Methods lab (https://therocinante-lab.github.io). His research focuses on cutting edge technologies encompassing molecular -omics, statistics and breeding to research strategies that contribute to the development of superior crop varieties. Our focus involves genomic prediction and selection, association mapping and characterization of allelic diversity

Dr. Marco Antônio Peixoto

Associated Researcher
Quantitative Genetics
UF/IFAS Horticulture Sciences
Sweet Corn Breeding and Genomics

Dr. Peixoto is a quantitative geneticist who specializes in bridging statistical and genetic methods to enhance plant breeding programs. He is dedicated to developing algorithms and using advanced techniques to optimize quantitative traits and improve breeding programs. More specifically, Dr. Peixoto is applying genomic methods to sweet corn hybrid prediction, as well as using simulations tools and real data to harness long-term genetic gains while circumventing the decrease of variance by controlling inbreeding rates via cross usefulness prediction and mating plan optimization

Dr. Joshua A. Sleper

Quantitative Genetics & Breeding
UF/IFAS Strawberry Breeding Program

Dr. Sleper is a quantitative genetics researcher in the Horticultural Sciences Department at the University of Florida. He has worked on the UF/IFAS strawberry breeding team since 2023 leading efforts to improve genomic selection capabilities and develop phenomics-aided breeding methods. Prior to his work at UF, he worked for a decade in industry, heading the development of a global genomic selection breeding pipeline for maize and leading a breeding technology team exploring new methods to optimize plant breeding processes. A major focus of his research is developing high-throughput methods to improve genetic gain and bringing his industry experience in maize breeding to the UF strawberry breeding program.

Martín Zapien

Ph.D. Candidate
Research Assistant
UF/IFAS Blueberry Breeding Program

Mr. Zapien, originally from Mexico, holds a bachelor's in Ag & Food Industries Engineering from El Tecnologico de Zamora. He earned a master's in Horticultural Sciences and a certificate in Biological Systems Modeling from the University of Florida/IFAS , where he is now a Ph.D. candidate. His research focuses on blueberry ripening and strategies to improve productivity and harvest efficiency, including developing genomic selection models for optimizing harvest operations of blueberry farms in Florida.

Detailed Agenda

Monday, July 7, 2025 (NVIDIA-Malachowsky)	
1889 Museum Rd, Gainesville, FL 32606	
7:30am-8:10am	Registration
8:10am-8:30am	Welcome and Introduction from Dr. John Davis, Associate Dean for Research for UF/IFAS and Dr. Jason Ferrell, Professor and Department Chair Agronomy Department
8:30am-9:50am	Dr. Jarquín, Ph.D Candidate Julian García-Abadillo, and Ph.D. Candidate Vitor Sagae, and Dr. Ying Zhang: Intro to R, CHiDO and HiPerGator.
9:50am-10:10am	Break (Group Photo)
10:10am-11:30am	Dr. Ríos - Field Plot Designs
11:30am-1:00pm	Lunch on your own
1:00pm-2:20pm	Dr. Resende & Dr. Peixoto - Intro to Genomic Selection (GS)- and Genomic Prediction (GP) Models (Bayesian Alphabet) & Application in Maize
2:20pm-2:40pm	Break
2:40pm-4:00pm	Dr. Pérez-Rodríguez: Bayesian Factor Analytic and Other Covariance Structures, multi-trait multi-environment strategies
6:00pm-9:00pm	Social Activities TBD

Tuesday, July 8, 2025 (NVIDIA-Malachowsky)	
1889 Museum Rd, Gainesville, FL 32606	
8:30am-9:50am	Dr. Jarquín, Ph.D Candidate Julian García-Abadillo, and Ph.D. Candidate Vitor Sagae: G×E Prediction Models Using Weather Data
9:50am-10:10am	Morning Break
10:10am-11:30am	Dr. Qian Zhao: Boost Data Science Workflows with GPU Acceleration
11:30am-1:00pm	Lunch Break on your own
1:00pm-2:20pm	Dr. Vieira: GS in Major Crops (Soybean)
2:20pm-2:40pm	Afternoon Break
2:40pm-4:00pm	Ph.D. Candidate Julian García-Abadillo: AI Methods for Genomic Prediction I
4:00pm-4:30pm	Dr. Sleper: Integrating phenomic and genomic data in the University of Florida strawberry breeding program
6:00pm-9:00pm	Social Activities TBD

Wednesday, July 9, 2025 (NVIDIA-Malachowsky)	
1889 Museum Rd, Gainesville, FL 32606	
8:30am-9:50am	Dr. Jarquín, Ph.D. Candidate Vitor Sagae, Ph.D. Candidate Julian García-Abadillo, Ph.D. Candidate Martín Zapien: strategies for large dimension datasets (large n and large p), Sparse Testing Designs, and prediction of production curves.
9:50am-10:10am	Morning Break
10:10am-11:30am	Dr. Julio Isidro y Sánchez: Rethinking Optimal Genomic Mating. A Closer Look at the Usefulness Criterion
11:30am-1:00pm	Lunch on your own
1:00pm-2:20pm	Dr. Hidalgo - Developments in ssGBLUP and ssGWAS I
2:20pm-2:40pm	Afternoon Break
2:40pm-4:00pm	Dr. Hidalgo: Developments in ssGBLUP and ssGWAS II
6:00pm-9:00pm	**Social Activities** TBD

Thursday, July 10, 2025 (NVIDIA-Malachowsky)	
1889 Museum Rd, Gainesville, FL 32606	
8:30am-9:50am	Dr. Jarquín and Ph.D. Candidate García-Abadillo - Multi-Omic Integration for Continuous and Categorical Data
9:50am-10:10am	Morning Break
10:10am-11:30am	Dr. Pérez-Rodríguez: Genomic Prediction Models for Predicting Hybrid Performance Using Genomic Information from Inbreds (GCA and SCA)
11:30am-1:00pm	Lunch on your own
1:00pm-2:20pm	Ph.D Candidate Julian García-Abadillo Velazco, and Dr. Jarquín: AI Methods for Genomic Prediction II / Deep Learning Optimization
2:20pm-2:40pm	Afternoon Break
2:40pm-4:00pm	Dr. Huiwen Ju - NVIDIA LLMs for genomics research and AI research agents
6:00pm-9:00pm	**Social Activities** TBD

Friday, July 11, 2025 (NVIDIA-Malachowsky)	
1889 Museum Rd, Gainesville, FL 32606	
8:30am-9:50am	Dr. Pérez-Rodríguez: Using Sparse Matrices in BGLR for Analysis of Field Experiments
9:50am-10:10am	Morning Break
10:10am-11:30am	Dr. Messina: DSSAT / Crop Growth models and AI
11:30am-1:00pm	Lunch Break
1:00pm-2:20pm	Dr. Enid Pérez-Lara: Women in Multi-Omic Integration and Applied AI Breeding - Breaking Barriers Through Data and Technology
2:20pm-2:40pm	Afternoon Break
2:40pm-4:00pm	Dr. Hsiao-Yi Hung: Implementing Predictive Breeding Pipeline with Publicly Available Software - Echidna
6:00pm-9:00pm	Social Activities TBD

PRESENTATIONS TIME

Session are listed in chronological order

Intro to R, CHiDO and HiPerGator

Monday, July 7, 2025 | 8:30AM - 9:50AM

Dr. Diego Jarquín

Mr. Julián García-Abadillo

Mr. Vitor Sagae

Dr. Ying Zhang

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Field Plot Designs

Monday, July 7, 2025 | 10:10AM - 11:30AM

Dr. Esteban Ríos

NOTES

Intro to Genomic Selection (GS)– and Genomic Prediction (GP) Models (Bayesian Alphabet) & Application in Maize

Monday, July 7, 2025 | 1:00PM - 2:20PM

Dr. Marcio Resende

Dr. Marco Peixoto

NOTES

Bayesian Factor Analytic and Other Covariance Structures, Multi-Trait Multi-Environment Strategies

Monday, July 7, 2025 | 2:40PM - 4:00PM

Dr. Paulino Pérez-Rodríguez

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G×E Prediction Models Using Weather Data

Tuesday, July 8, 2025 | 8:30AM - 9:50AM

Mr. Vitor Sagae

Dr. Diego Jarquín

Mr. Julián García-Abadillo

NOTES

Boost Data Science Workflows with GPU Acceleration

Tuesday, July 8, 2025 | 10:10AM - 11:30AM

Dr. Qian Zhao

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GS in Major Crops (Soybean)

Tuesday, July 8, 2025 | 1:00PM - 2:20PM

Dr. Caio Canella Vieira

NOTES

AI Methods for Genomic Prediction I

Tuesday, July 8, 2025 | 2:40PM - 4:00PM

Mr. Julián García-Abadillo

NOTES

Integrating Phenomic and Genomic Data in the University of Florida Strawberry Breeding Program

Tuesday, July 8, 2025 | 4:00PM - 4:30PM

Dr. Joshua A. Sleper

NOTES

Strategies for Large Dimension Datasets (large n and large p), Sparse Testing Designs, and Prediction of Production Curves.

Wednesday, July 9, 2025 | 8:30AM - 9:50AM

Dr. Diego Jarquín

Mr. Julián García-Abadillo

Mr. Vitor Sagae

Mr. Martín Zapien

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Rethinking Optimal Genomic Mating. A Closer Look at the Usefulness Criterion

Wednesday, July 9, 2025 | 10:10AM - 11:30AM

Dr. Julio Isidro y Sánchez

NOTES

Developments in ssGBLUP and ssGWAS I

Wednesday, July 9, 2025 | 1:00PM - 2:20PM

Dr. Jorge Hidalgo

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Developments in ssGBLUP and ssGWAS II

Wednesday, July 9, 2025 | 2:40PM - 4:00PM

Dr. Jorge Hidalgo

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Multi-Omic Integration for Continuous and Categorical Data

Thursday, July 10, 2025 | 8:30AM - 9:50AM

Dr. Diego Jarquín

Mr. Julián García-Abadillo

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Genomic Prediction Models for Predicting Hybrid Performance Using Genomic Information from Inbreds (GCA and SCA)

Thursday, July 10, 2025 | 10:10AM - 11:30AM

Dr. Paulino Pérez-Rodríguez

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AI Methods for Genomic Prediction II/ Deep Learning Optimization

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Mr. Julián García-Abadillo

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NVIDIA LLMs for Genomics Research and AI Research Agents

Thursday, July 10, 2025 | 2:40PM - 4:00PM

Dr. Huiwen Ju

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Using Sparse Matrices in BGLR for Analysis of Field Experiments

Friday, July 11, 2025 | 8:30AM - 9:50AM

Dr. Paulino Pérez-Rodríguez

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DSSAT / Crop Growth Models and AI

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Dr. Charlie Messina

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Women in Multi-Omic Integration and Applied AI Breeding – Breaking Barriers Through Data and Technology

Friday, July 11, 2025 | 1:00PM - 2:20PM

Dr. Enid Pérez-Lara

NOTES

Implementing Predictive Breeding Pipeline with Publicly Available Software – Echidna

Friday, July 11, 2025 | 2:40PM - 4:00PM

Dr. Hsiao-Yi Hung

NOTES

ADDITIONAL INFORMATION

Internet Access

To access Internet in the meeting space, follow these instructions on your device:
1. Connect to the network “ufguest”
2. You will be directed to the UF Guest Network Acceptable Use Policy.
3. Click on "Accept AUP and Connect"

Name Badge

Your name badge serves as your admission to all functions while attending the short course.

Breaks & Lunches

Mid-morning and afternoon breaks provide an opportunity to stretch legs and use bathroom. A full, 90-minute lunchtime on your own on Monday through Friday at the Reitz Union.

Cell Phones, Mobile Devices, Tablet Devices

Please mute your cell phones, laptops, tablets, and mobile devices while in the meeting room. Please respect the instructors’ wishes not to share certain sensitive data on social media. Please do not photograph or share on social media.

Lost & Found

If you lose or find an article, check with the onsite registration staff.

Code of Conduct

All meeting participants must agree to follow our Code of Conduct when they register. We welcome you to join, sustain, foster, and help grow our inclusive and supportive environment.

Medical Emergency and Safety Information:

In the event you experience any type of emergency, please dial 911 to summon medical assistance or to contact authorities.

Thank you Sponsors!

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Special Thanks to: our Volunteers

Edmundo R. Caballero Espinosa

Samuel Flores Galarza

Juan David Jimenez Pardo

Sehijpreet Kaur

Shatabdi D. Proma

Xiangming Zhou


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