Paulo Izquierdo

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EDUCATION

- 2023 Ph.D; Plant Breeding, Genetics, and Biotechnology, Department of Plant, Soil, and Microbial Sciences, Michigan State University, East Lansing, MI.
- 2022 Graduate certificate, Computational Plant Science, Department of Plant Biology, Michigan State University, East Lansing, MI.
- 2010 B. S; Biology, Universidad del Tolima, Ibagué, Colombia.

RESEARCH AND PROFESSIONAL EXPERIENCE

RESEARCH AND I ROFESSIONAL EXIENCE	
03/2023-Present	Postdoctoral Research Associate. Investigate the genetic architecture of complex traits in plant species and incorporate these findings into genomic prediction models. Great Lakes Bioenergy Research Center. Principal Supervisor: Shin-Han Shiu
01/2017-12/2022	Graduate Research Assistant. Exploring the genetic architecture and improving genomic prediction accuracy for yield and end-use quality traits in common bean. Michigan State University. Principal Supervisor: Karen Cichy
06/2022-08/2022	Quantitative Genetics / Computational Biology Intern. Identified potential targets for genome editing using machine learning. INARI.
06/2016-12/2016	Visiting Scholar. Developed molecular markers linked to anthracnose resistance in common bean. Michigan State University. Principal Supervisor: Karen Cichy and James Kelly
09/2013-06/2016	Research Assistant. Uncovered the genetic architecture of the mineral concentration in common bean using GWAS and QTL approaches in a Multiparent Advance Generation Intercross population. CIAT. Principal Supervisor: Bodo Raatz
01/2011-08/2013	Research Assistant. Identified genomic regions associated with agronomic traits in sugarcane. Colombian Sugarcane Research Center (Cenicaña). Principal Supervisor: Jershon Lopez
07/2010-12/2010	Visiting Researcher. Implemented marker-assisted selection for resistance to anthracnose in common bean. CIAT/Universidad Nacional de Colombia. Principal Supervisor: Matthew Blair
01/2009-06/2010	Undergraduate Research Assistant. Evaluated the use of wild germplasm to increase the mineral concentration of cultivars in common beans. CIAT. Principal Supervisor:

PUBLICATIONS

https://scholar.google.com/citations?user=5sZkDtAAAAAJ&hl=en

Matthew Blair

Peer-Reviewed Journal Articles:

- **Izquierdo P**, Sadohara R, Wiesinger J, Glahn R, Urrea C, Cichy K. Genome-wide association and genomic prediction for iron and zinc concentration and iron bioavailability in a collection of yellow dry beans. Front. Genet. 15, 1-12. https://doi.org/10.3389/fgene.2024.1330361
- **Izquierdo P**, Kelly J, Beebe S, Cichy K. Combination of meta-analysis of QTL and GWAS to uncover the genetic architecture of seed yield and seed yield components in common bean. The Plant Genome, 16, e20328. https://doi.org/10.1002/tpg2.20328

- Amongi W, Nkalubo S, Ochwo-Ssemakula M, Badji A, Dramadri I, Odongo T, Nuwamanya E, Tukamuhabwe P, **Izquierdo P**, Cichy K, Kelly J, Mukankusi C. Phenotype based clustering, and diversity of common bean genotypes in seed iron concentration and cooking time. PLOS ONE 18(5): e0284976. https://doi.org/10.1371/journal.pone.0284976
- Amongi W, Nkalubo S, Ochwo-ssemakula M, Dramadri I, Odongo T, Nuwamanya E, Tukamuhabwe P, **Izquierdo P**, Cichy K, Kelly J, Mukankusi C. Genetic clustering, and diversity of African panel of released common bean genotypes and breeding lines. Genet. Resour. Crop Evol. 70, 2063–2076. https://doi.org/10.1007/s10722-023-01559-y
- 2022 Sadohara R, **Izquierdo P**, Couto Alves F, Porch T, Beaver J, Urrea C, Cichy K. The *Phaseolus vulgaris* L. Yellow Bean Collection: genetic diversity and characterization for cooking time. Genet. Resour. Crop. Evol. 69, 1627–1648. https://doi.org/10.1007/s10722-021-01323-0
- Sadohara R, Long Y, **Izquierdo P**, Urrea C, Morris D, Cichy K. Seed coat color genetics and genotype × environment effects in yellow beans via machine-learning and genome-wide association. Plant Genome 15:e20173. https://doi.org/10.1002/tpg2.20173
- Diaz S, Ariza-Suarez D, **Izquierdo P**, Lobaton J, de la Hoz J, Acevedo F, Duitama J, Guerrero J, Cajiao C, Mayor V, Beebe S, Raatz B. Genetic mapping for agronomic traits in a MAGIC population of common bean (*Phaseolus vulgaris* L.) under drought conditions. BMC Genomics 21, 799. https://doi.org/10.1186/s12864-020-07213-6
- Berry M, **Izquierdo P**, Jeffery H, Shaw S, Nchimbi-Msolla S, Cichy K. QTL analysis of cooking time and quality traits in dry bean (*Phaseolus vulgaris* L.). Theoretical and Applied Genetics 133, 2291–2305. https://doi.org/10.1007/s00122-020-03598-w
- 2018 **Izquierdo P**, Astudillo C, Blair M, Iqbal A, Raatz B, Cichy K. Meta-QTL analysis of seed iron and zinc concentration and content in common bean (*Phaseolus vulgaris* L.). Theoretical and Applied Genetics 131, 1645–1658. https://doi.org/10.1007/s00122-018-3104-8
- **Izquierdo P**, Shaw S, Berry M, Cichy K. A saturated genetic linkage map of common bean (*Phaseolus vulgaris* L.) developed using Genotyping by Sequencing (GBS). Annual report of the Bean Improvement Cooperative 60.
- Perea C, De La Hoz J, Cruz D, Lobaton J, **Izquierdo P**, Quintero J, Raatz B, Duitama J. Bioinformatic analysis of genotype by sequencing (GBS) data with NGSEP. BMC Genomics 17, 498. https://doi.org/10.1186/s12864-016-2827-7
- Blair M, **Izquierdo P**, Astudillo C, Grusak M. A legume biofortification quandary: Variability and genetic control of seed coat micronutrient accumulation in common beans. Front. Plant Sci. 4, 1–14. https://doi.org/10.3389/fpls.2013.00275
- Delgado H, Pinzón E, Blair M, **Izquierdo P**. Evaluation of bean (*Phaseolus vulgaris* L.) lines result of an advanced backcross between a wild accession and Radical Cerinza. U.D.C.A Act. & Div. Cient. 16(1), 79–86. https://doi.org/10.31910/rudca.v16.n1.2013.861
- 2013 **Izquierdo P**, Gutierrez A, Victoria J, Angel J, Avellaneda M, Lopez J. Molecular markers associated with resistance to Sugarcane yellow leaf virus. International Society of Sugar Cane Technologists 28, 1179–1188.
- Blair M, **Izquierdo P**. Use of the advanced backcross-QTL method to transfer seed mineral accumulation nutrition traits from wild to Andean cultivated common beans. Theoretical and Applied Genetics 125, 1015–1031. https://doi.org/10.1007/s00122-012-1891
- Blair M, **Izquierdo P**, Astudillo C, Monserrate F, Cortés M, Avila P, Felde T, Pfieffer W. Utilization of near infrared spectrophotometry (NIRS) analysis for evaluation of mineral content in Andean bean samples. Annual report of the Bean Improvement Cooperative 57.

Pre-prints:

Palande S, Arsenault J, Basurto P, Bleich A, Brown B, Buysse S, Connors N, Adhikari S, Dobson K, Guerra F, Guerrero M, Harlow S, Herrera H, Hightower A, **Izquierdo P**, et al. A data-driven evaluation of Arabidopsis-centric research and the model species concept. bioRxiv. https://doi.org/10.1101/2023.08.20.554029

In preparation:

Izquierdo P, Kelly J, Cichy K. Accelerating genetic gain in dry beans for yield and end-use quality traits using genomic selection and high-throughput phenotyping.

SCIENTIFIC PRESENTATIONS

Invited talks:

- Nutritional Quality: An Essential Trait for New Crop Varieties, presented during the Webinar "Agriculture, Biodiversity, and Nutrition: Foundations for Sovereign, Sustainable, and Resilient Food Systems". Universidad Nacional de Colombia. April 22.
- Exploring the genetic architecture and improving genomic prediction accuracy for yield, mineral concentration, and canning quality traits in dry bean (*Phaseolus vulgaris* L.). presented during the Webinar "Phytopathology and Plant Breeding". Universidad de Nariño, Colombia. November 3.
- 2020 Graduate Studies: Do's and Don'ts, presented at the "Caribbean Microbial Meeting". October 29.
- 2020 Use of molecular markers in dry bean breeding, Universidad Industrial de Santander, Colombia. June 24.

Contributed talks and posters:

- **Izquierdo, P**; Segura, K; Weng, X; Juenger, T; Lowry, D and Shiu, S. Unraveling the genetic landscape of biomass in switchgrass through transcriptomics. Plant Biology. **Poster.**
- 2024 **Izquierdo, P**; Segura, K; Weng, X; Juenger, T; Lowry, D and Shiu, S. Uncover the genetic architecture of agronomic traits in a switchgrass diversity panel across two environmentally distinct locations: Michigan and Texas. Great Lakes Bioenergy Research Center Annual Science Meeting. **Oral presentation.**
- 2019 **Izquierdo, P**; Lopez, M; Kelly, J; Cichy, K. Assessing Genomic Selection Prediction Accuracy for Yield and End-Use Quality Traits in Black Beans. Bean improvement cooperative. **Poster.**
- 2019 **Izquierdo, P**; Katuuramu, D; Cichy, K. Genomic selection for nutritional traits and cooking time in common bean) using Genotyping by Sequencing. Plant & Animal Genome. **Poster.**
- 2018 **Izquierdo, P**; Astudillo, C; Iqbal, A; Blair, M; Raatz, B and Cichy, K. Meta-QTL Analysis in Common Bean to Uncover the Genetic Architecture of Iron and Zinc Concentration in Seed. Plant & Animal Genome. **Poster.**
- **Izquierdo, P**; Shaw, S; Berry, M and Cichy K. A saturated genetic linkage map of common bean developed using Genotyping by Sequencing (GBS). Bean improvement cooperative. **Poster.**
- 2016 **Izquierdo, P**; Lobaton, J; Mayor, V; Grajales, M; Cajiao, C; Duitama, J and Raatz, B. Genomewide association mapping for yield and other agronomic traits in a Multi-parent advanced generation inter-cross population of Mesoamerican common bean (*Phaseolus vulgaris L.*). IX Latin American and Caribbean Agricultural and Forestry Biotechnology Meeting, Peru. **Oral presentation.**
- **Izquierdo, P**; Gutiérrez, A; Victoria, J; C; Ángel, López, J; Avellaneda, C. Molecular markers associated with resistance to the sugarcane yellow leaf virus. XXVIII International Society of Sugarcane Technologists (ISSCT). Brazil. **Oral presentation.**

- 2012 **Izquierdo, P**; Gutiérrez, A; Victoria, J; Ángel, J; López, J; Avellaneda, C. Molecular markers associated with resistance to the sugarcane yellow leaf virus. IX Association for Sugarcane Technology in Latin America and the Caribbean (Atalac-Tecnicaña). Colombia. **Oral presentation.**
- 2011 **Izquierdo, P**; Gutiérrez, A; Avellaneda, C; Victoria, J; Ángel, López, J. Molecular markers associated with resistance to the sugarcane yellow leaf virus. Colombian and Latin American Phytopathological Association. Colombia. **Oral presentation.**

HONORS, AWARDS, AND FUNDING

- Norman and Jessie Thompson fellowship in Crop and Soil Sciences (\$5,000).
- NSF Research Traineeship Integrated training model in plant and computational sciences fellowship (\$29,781).
- 2021 Everett and Jane Everson fellowship in Plant Breeding (\$2,500).
- 2021 Jason and Dana Lilly fellowship in Plant Breeding, Genetics & Biotechnology (\$1,500).
- Norman and Jessie Thompson fellowship in Crop and Soil Sciences (\$4,000).
- 2021 College of Agriculture and Natural Resource fellowship, Michigan State University (\$6,000).
- 2020 Bayer Diversity Initiative Scholar.
- 2019 Everett and Jane Everson fellowship in Plant Breeding (\$2,500).
- 2019 Norman and Jessie Thompson fellowship in Crop and Soil Sciences (\$1,000).
- 2019 Graduate student language fellowship in undergraduate teaching and learning Residential College in the Arts and Humanities, Michigan State University (\$4,000).
- 2019 Council of Graduate Students, Conference Award, Michigan State University (\$300).
- 2018 The Crop and Soil Science Graduate award, Michigan State University (\$1,700).
- 2018 Jason and Dana Lilly fellowship in Plant Breeding, Genetics & Biotechnology (\$1,500).
- 2018 Elmer C. Rossman fellowship in Plant, Soil & Microbial Sciences (\$3,500).
- 2018 Everett and Jane Everson fellowship in Plant Breeding (\$2,500).
- 2018 Resilient and Nutritious Dry Beans for Africa fellowship, USDA-FAS (\$5,400).
- 2017 Doctoral Fellowship Program. COLCIENCIAS, Colombia's Administrative Department of Science, Technology, and Innovation (\$120,000).

TEACHING EXPERIENCE

Michigan State University:

- Teaching Assistant: Department of Plant, Soil and Microbial Sciences, Introduction to Plant Genetics (CSS350), Michigan State University
- 2019 Spanish Language Assistant: Residential College in the Arts and Humanities, Program on Sustainability in Costa Rica, Michigan State University.

Workshops:

- 2024 "Introduction to Machine Learning", Great Lakes Bioenergy Research Center Annual Science Meeting. Organizer and lecturer for 1-session workshop, 60 participants.
- 2023 "Introduction to Machine Learning", Universidad Industrial de Santander, Colombia. Organizer and lecturer for 2-session workshop, 32 participants.
- 2021 "Quantitative Genetics", MSU-Feed the Future Innovation Lab for Crop Improvement (Eastern Africa). Organizer and lecturer for 9-session workshop, 123 participants. https://pauloizquierdo.github.io/Quantitative_Genetics/

- 2021 "Quantitative Genetics", Universidad Industrial de Santander, Colombia. Organizer and lecturer for 1-session workshop, 52 participants. https://compasscol.github.io/2021B_talleres-UIS.html
- "Data visualization with R", Universidad Industrial de Santander, Colombia. Organizer and lecturer for 2-day workshop, 40 participants. https://compasscol.github.io/dataviz/
- 2021 "Introduction to R", Universidad Industrial de Santander, Colombia. Organizer and lecturer for 2-day workshop, 40 participants. https://compasscol.github.io/IntroR/

MENTORING

- Yuranis Miranda, Graduate Research Assistant, Universidad Industrial de Santander, Colombia. Project: Genetic and Morphological Variability of *Vaccinium meridionale* Sw. Across the Central and Eastern Andean Regions of Colombia.
- Anisa Rashid, Undergrad researcher/hourly worker, Michigan State University. Project: Evaluation of canning quality in dry beans.
- Winnyfred Amongi, Visiting Scholar, Michigan State University. Project: High-Throughput genotyping and genomic prediction in dry beans. *co-authored paper*.
- Wilson Santiago, Biology Intern. International Center for Tropical Agriculture. Project: Genetic mapping for agronomic traits in a MAGIC population of common bean. *co-authored paper*.
- 2016 Laura Paz, Biology Intern. International Center for Tropical Agriculture. Project: Fine mapping of regions associated with Fe concentration in dry bean.

STUDENT MENTORING COMMITTEES

- 2023 Miguel Mendoza, Ph.D. Universidad Nacional de Colombia, Bogotá, Colombia. Project: Genetics of Nitrogen Use Efficiency in Plants.
- Hernán Maigual, MSc. Universidad de Nariño, Pasto, Colombia. Project: Variability Assessment of Fava Beans (*Vicia faba* L.) Originating from the Andean Region of Nariño, Colombia.
- Diego González, MSc. Universidad Nacional de Colombia, Bogotá, Colombia. Project: Analysis of the Fungal Microbiome in Cocoa Soils with Varying Cadmium Concentrations.
- 2021 Emili Garcia, MSc. Universidad Nacional de Colombia, Bogotá, Colombia. Project: Identification of Allelic Variants and Assessment of Candidate Gene Expression Linked to Bacterial Vascular Wilt Resistance in Cassava.
- 2020 Fabián Villamil, MSc. Universidad Nacional de Colombia, Bogotá, Colombia. Project: Expression of the polyhydroxyalkanoate (PHA) synthase enzyme from Aeromonas caviae in transgenic tobacco, lead to the synthesis of a poly (4-hydroxybutyrate)

PROFESSIONAL SERVICE

Ad Hoc Grant Reviewer:

2023 Colombia's Administrative Department of Science, Technology, and Innovation (Colciencias).

Ad Hoc Reviewer:

Scientific reports (1), GigaScience (2), Agronomy (1), PhytoFrontiers (1).

Conference/symposium organization:

- Co-Organizer, Workshops series focused on data analysis in biology and agriculture. Universidad Industrial de Santander, Colombia. https://compasscol.github.io/2021B_talleres-UIS.html
- 2021 Co-Organizer, Online seminar series on phytopathology and plant breeding. Universidad de Nariño, Colombia. https://compasscol.github.io/2021A_Conferencias-UNar.html

- 2020 Co-Founder, Community Platform for Agricultural Sciences (COMPASS). Website: https://compasscol.github.io/
- 2020 Co-Organizer, Ciencia en línea: impulsando el uso de plataformas para la divulgación, colaboración e inspiración científica, symposium, SOCOLEN
- 2019 Co-Organizer, Phenomic Application in Plant Breeding, Corteva-PBGB symposium.

MEMBERSHIPS

American Society of Plant Biologists