Rodrigo Rampazo Amadeu

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Education

Expected August 2021 Pн.D. in Horticultural Sciences (Minor: Statistics), University of Florida, USA Dissertation: "Statistical methods for genomic-assisted blueberry breeding"

Advisor Dr. Patricio Munoz, Blueberry Breeding & Genomics Lab

2018

M.S. in Plant Genetics and Breeding, ESALQ, University of São Paulo, Brazil Thesis: "Molecular pairwise relatedness in autopolyploids: a simulation study"

Advisor Dr. Antonio Augusto Franco Garcia, Statistical-Genetics Lab

B.Eng. in Agriculture (Minor: Biotechnology), ESALQ, University of São Paulo, Brazil

B.Edu. in Agricultural Sciences, ESALQ, University of São Paulo, Brazil

Research Experience

2018-present

University of Florida, USA

Graduate Research Assistant, Blueberry Breeding & Genomics Lab, Supervisor Dr. Patricio Munoz

- Planning and optimization of breeding program
- Genetic data analysis (genomic prediction, population genetics, gene discovery and mapping)
- Agricultural data analysis (experiment design and planning, linear mixed models)
- Development of statistical-genetics software (AGHmatrix, diaQTL, PedigreeSimR)
- Plant breeding activities (phenotyping, selection)

2016-2018

University of São Paulo, Brazil

Graduate Research Assistant, Statistical-Genetics Lab, Supervisor Dr. Augusto F. Garcia

- Genetic data analysis (genomic prediction, population genetics, gene discovery and mapping)
- Agricultural data analysis (experiment design and planning, linear mixed models)
- Development of statistical-genetics software (onemap, onemap2pop, fullsibQTL)
- CNPq Scholarship

2014

University of Florida, USA

Intern, Forage Breeding and Genomics Lab, Supervisor Dr. Patricio Munoz

- Development of software to build genomic relationship matrices (AGHmatrix)
- Plant breeding activities (pollination, phenotyping, selection)
- Science without Borders Scholarship

2010-2015

University of São Paulo, Brazil

Intern, Statistical-Genetics Lab, Supervisor Dr. Augusto F. Garcia

- Development of pipeline for SNP dosage calling in autopolyploid data
- Population structure analysis of sugarcane panel
- CNPq & Santander scholarships

Awards & Scholarships

2020	Scholarship, Murial Rumsey scholarship, CALS, Univ. of Florida
2020	Scholarship, outstanding teaching assistantship, Univ. of Florida
2019	Scholarship, outstanding teaching assistantship, Univ. of Florida
2019	Award, poster competition Plant Science Symposium, Univ. of Florida - 1st Place
2016	Award, Prof Brieger, best graduating student of Dep. of Genetics, Univ. of São Paulo
2013	Scholarship, Science without Borders - CAPES - 1yr at Univ. of Florida
2012	Scholarship, Scientific Initiation - PIBIC/CNPq - 1yr
2011	Scholarship, Scientific Initiation - Santander - 1yr

Journal Articles

- SUBMITTED MANUSCRIPTS (3)
- Amadeu, RR; Munoz, R; Zheng, C; Endelman, JB. "QTL mapping in outbred tetraploid (and diploid) diallel populations". *bioRxiv*, link
- Zheng, C; **Amadeu, RR**; Munoz, R; Endelman, JB. "Haplotype reconstruction in connected tetraploid F1 populations". *bioRxiv*, link
- Gazaffi, R*; **Amadeu, RR*** (*contributed equally for this study); Mollinari, M; Rosa, JRBF; Taniguti, CH; Margarido, GRA; Garcia, AAF. "fullsibQTL: an R package for QTL mapping in biparental populations of outcrossing species". bioRxiv, link
 - Published in Peer-Reviewed Journals (15)
- Ferrao, LFV; **Amadeu, RR**; Benevenuto, J; de Bem Oliveira, I; Munoz, R. "Genomic prediction in an outcrossing and autotetraploid fruit crop: lessons from blueberry breeding". *Front. Plant Sci.*, TBA, link
- Quezada, M; **Amadeu, RR**; Vignale, B; Cabrera, D; Pritsch, C; Garcia, AAF. "Construction of a high-density genetic map of *Acca sellowiana* (Berg.) Burret, an outcrossing species, based on two connected mapping populations". *Front. Plant Sci.*, 12 (626811), link
- Cappai, F*; **Amadeu, RR*** (*contributed equally for this study); Benevenuto, J; Cullen, R; Garcia, AL; Grossman, AY; Ferrão, LFV; Munoz, PR. "High-resolution linkage map and QTL analyses of fruit firmness in autotetraploid blueberry". Front. Plant Sci., 11 (562171), link
- de Bem Oliveira, I; **Amadeu, RR**; Ferrão, LFV; Munoz, PR. "Optimizing whole-genomic prediction for autotetraploid blueberry breeding". *Heredity*, 125, link
- Amadeu, RR; Lara, LADC; Munoz, PR; Garcia, AAF. "Estimation of molecular pairwise relatedness in autopolyploid crops". *G*₃, 10(12), link
- de Oliveira, AA; Resende, MFR; Ferrão, LFV; **Amadeu, RR**; Guimarães, LJM; Guimarães, CT; Pastina, MM; Margarido, GRA. "Genomic prediction applied to multiple traits and environments in second season maize hybrids". *Heredity*, 125, link
- Amadeu, RR; Ferrão, LFV; de Bem Oliveira, I; Benevenuto, J; Endelman, JB; Munoz, PR. "Impact of dominance effects on autotetraploid genomic prediction". *Crop Science*, 60(2), link
- Estrada-Reyes, ZM; Tsukahara, Y; **Amadeu, RR**; Goetsch, AL; Gipson, TA; Sahlu, T; Puchala, R; Wang, Z; Hart, ST; Mateescu, RG. "Signatures of selection for resistance to *Haemonchus contortus* in sheep and goats". *BMC Genomics*, 20(1), link
- Lara, LADC; Santos, MF; Jank, L; Chiari, L; Vilela, MDM; **Amadeu, RR**; dos Santos, JP; Pereira, GDS; Zeng, ZB; Garcia, AAF. "Genomic selection with allele dosage in *Panicum maximum* Jacq.". *G*₃, 9(8) link
- Benevenuto, J; Ferrão, LFV; **Amadeu, RR**; Munoz, P. "How can a high-quality genome assembly help plant breeders?". *GigaScience*, 8(6), link
- de Bem Oliveira, I; Resende Jr, MFR; Ferrao, LFV; **Amadeu, RR**; Endelman, JB; Kirst, M; Coelho, ASG; Munoz, PR. "Genomic prediction of autotetraploids; influence of relationship matrices, allele Dosage, and continuous genotyping calls in phenotype prediction". *G*₃, 9(4), link
- Conson, ARO*; Taniguti, CH*; **Amadeu, RR*** (*contributed equally for this study); Andreotti, IAA; de Souza, LM; dos Santos, LHB; Rosa, JRBF; Mantello, CC; da Silva, CC; Scaloppi Jr, EJ; Ribeiro, RV; Le Guen, V; Garcia, AAF; Gonçalves, PS; Souza, AP. "High-resolution genetic map and QTL analysis of growth-related traits of Hevea brasiliensis". Front. Plant Sci., 9(1255), link
- Ferreira, DA; Abreu, GF; Cheavegatti-Gianotto, A; Soldi, MCM; Carneiro, MS; **Amadeu, RR**; Hoffmann, HP; Aricetti, JA; Wolf, LD; Caldana, C. "Metabolite profiles of sugarcane culm reveal the relationship among metabolism and axillary bud outgrowth in genetically related sugarcane commercial cultivars". *Front. Plant Sci.*, 9(857), link
- Cellon, C; **Amadeu, RR**; Olmstead, JW; Mattia, MR; Ferrao, LFV; Munoz, PR. "Estimation of genetic parameters and prediction of breeding values in an autotetraploid blueberry breeding population with extensive pedigree data". *Euphytica*, 214(87), link

2016

Amadeu, RR; Cellon, C; Olmstead, JW; Garcia, AAF; Resende, MF; Munoz, PR. "AGHmatrix: R package to construct relationship matrices for autotetraploid and diploid species, a blueberry example". *The Plant Genome*, 9(3), link

Selected Teaching and Talks Experience

Seminar "QTL mapping in tetraploid diallel populations". CGDG. The Roslin Institute, UK. 2021 **Seminar** "QTL mapping in autotetraploid multi-parent populations". The 6th International Confer-2020 ence of Quantitative Genetics (ICQG6). Virtual conference. link Teaching assistant of Field Plot Techniques, graduate level, University of Florida Fall 2020 Guest lecturer of Special Topic in Genetics and Breeding, graduate level, University of São Paulo Sum 2020 Fall 2010 Teaching assistant of Mol. Marker Assisted Plant Breeding, graduate level, University of Florida Teaching assistant of Field Plot Techniques, graduate level, University of Florida Fall 2019 Teaching assistant of Field Plot Techniques, graduate level, University of Florida Fall 2018 Teaching assistant of Calculus I, undergraduate level, University of São Paulo Spring 2015 **Teaching assistant** of Genetics, undergraduate level, University of São Paulo Spring 2012 Instructor of Precalculus in a college preparatory school, Piracicaba, Brazil 2011-2015

Ad-hoc Reviewer

Journals: BMC Genomics, Briefings in Bioinformatics

Software Development

AGHmatrix onemap author, compute relationship matrices for diploid and autopolyploid species, link contributor, build genetic maps in experimental crosses, link author, onemap extension to build multi-family genetic maps in outcrossing species, link co-author, QTL mapping in outcrossing species using composite interval mapping, link co-author, QTL mapping in multiparent and autopolyploid populations, link author, simulation of multiparent and autopolyploid populations, link

Skills

genetics genomic prediction, GWAS, QTL mapping, quantitative genetics, polyploid genetics, simulation statistics analysis of genetic & agricultural data, machine learning, linear mixed models R (advanced): package development, tidyverse, shiny/plotly app, parallelization shell/bash script, Linux, C/C++, ASReml, GitHub, AlphaGenes software, LTEX Portuguese (native), English (high proficiency)

Specialized Training

2021	Fundamentals Deep LearningDeep Learning Institute-NVIDIA, Gainesville, FL, US
2019	Analysis of Experiments Using ASReml-R, Dr. S. Gezan (UF), Gainesville, FL, US
2017	Modelling GxE Interaction in Genomic Prediction Analysis, Dr. J. Crossa (CIMMYT), Piracicaba
2016	Quantitative Genetics and Genomics Workshop, Drs. Morota & Spangler (UNL), Piracicaba, BR
2015	Brazilian Edition of the "Tucson Plant Breeding Institute", Org. Dr. Bruce Walsh, Piracicaba, BR
2015	EMBL-EBI Bioinformatics Workshop, EMBL-EBI, Piracicaba, BR