Department of Evolution and Ecology Center for Population Biology Genome Center University of California Davis

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Education

PhD Genetics, University of Georgia 2006 MS Botany, University of California Riverside 2000 BA Botany, University of California Riverside 1998

Academic Employment

Professor, Dept. Evolution and Ecology, University of California Davis 2019-present Professor, Dept. Plant Sciences, University of California Davis 2016-2019
Associate Professor, Dept. Plant Sciences, University of California Davis 2012-2016
Assistant Professor, Dept. Plant Sciences, University of California Davis 2009-2012
Postdoctoral Researcher, University of California Irvine 2006-2008
Profesor de Asignatura, Universidad Nacional Autónoma de México 2001

Selected Fellowships and Awards

Corn Pun Trophy, Genetics Society of America 2017
Stadler Mid-Career Excellence in Maize Genetics Award 2016
Faculty Development Award in recognition of university service 2015
DuPont Young Professor Award 2012
Presidential Early Career Award for Scientists and Engineers 2009
Dean's Award for Postdoctoral Excellence, UC Irvine 2008

Instruction and Advising

Current (total) advisees: 4 (22) postdoc, 5 (9) graduate, 3 (30) undergraduate Plant Biology (UC Davis, PLB200A, graduate), 2018-present Genetics (UC Davis, BIS 101, undergraduate), 2013-present Ecological Genomics (UC Davis, ECL 243, graduate), 2014-present Faculty advisor, US-Mexico graduate student exchange program, 2011-2015 Population and Quantitative Genetics (GGG 201D, graduate), 2010-2013 Plant Genetics (PLS 152, undergraduate), 2010-2011 Biología de Plantas I (undergraduate), UNAM, 2001

Service: selected from last 2 years

University

Chair, Ecology & Evolution Seminar Series

Campus-wide High-Performance Computing Task Force

Executive Cmte, Plant Biology Graduate Group

Ecology Graduate Group admissions committee

Campus Disciplinary Peer Review Committee on sexual violence and sexual harassment

Campus Task Force on Bioinformatics

Section Chair for Agricultural Plant Biology

Plant Sciences executive committee

Faculty advisor, Corteva graduate student symposium in plant science

Search committees: Climate Adaptation

Dept. of Plant Sciences academic planning committee

Professional

Chair, PEQG 2022 Conference

Organizing Committee, PEQG 2020 Conference

LEAD21 Leadership in Land Grant Institutions, Class 14

Skype-a-Scientist K-12 Outreach: Canada, Spain, New Jersey

Maize Genetics conference steering committee

Maize Genetics Awards Committee

Editorial Boards: Genes, Genomes, and Genetics (SE, AE)Genetics (Guest) PLoS Genetics (AE)PeerJ (SE, AE)PNAS (Guest)

Journal peer review: eLife (3), Cell, Nature Communications (3), Nature Reviews Genetics, PNAS (3), Science, PLoS Genetics, New Phytologist (2), Molecular Ecology, G3 (many), Plant Journal, Agriculture, Ecosystems and Environment

Current Funding

NSF: "Uncovering the role of transposons in maize variation" \$800,000 of \$4.5M total (Co-PI), 2019-2022

NSF: "Harnessing convergence and constraint to predict adaptations to abiotic stress for maize and sorghum" $$740,000 { of } $5.4M { total } (Co-PI), 2018-2022$

NSF: "The evolutionary genetics of pollen-pistil incompatibility and reproductive isolation in *Zea mays*" \$320,000 of \$1M total (Co-PI), 2018-2021

NSF: "The genetics of highland adaptation in maize", \$4.2M (PI), 2016-2021

Invited Seminars

Calvin Sperling Memorial Biodiversity Lectureship, Phoenix, 2020 NCSU, Raleigh, Nov 2020 U. Oulu, Finland, April 2020

NYU Abu Dhabi, UAE, Feb. 2020

U. Oregon, Eugene, Feb 2020

U. Massachusetts, Nov 2019

CAAS, Beijing, China, June 2019

U. of Science and Technology, Beijing, China, June 2019

International Forum on Crop Science, Wuhan, China, June 2019

U. California, Riverside, May 2019

U. Vermont, Burlington, Apr 2019

U. Washington, Seattle, Mar 2019

Keynote Speaker, Seed Central, U. California, Davis, Feb 2019

Annual James Brewbaker Lecture on Genetics & Plant Breeding, U. Hawaii, Manoa Dec. 2018

National Science Foundation, Washington DC, Sept 2018

U. Georgia, Athens, Aug 2018

Science and Society public lecture, Woodland, Aug 2018

Fisher Biosciences, July 2018

Texcoco pioneer symposium Sept. 2018 (declined)

Corteva Agriscience, Johnston, July 2018

Plenary Speaker, Maize Genetics Conference, Saint-Malo, France, Mar 2018

University of Paris-Saclay, Gif-sur-Yvette, France, Mar 2018

U. California, Davis, Mar 2018

Plant And Animal Genome Conference (two workshops), San Diego, Jan 2018

Plant And Animal Genome Conference (big data) (declined), Jan 2018

U. Nebraska, Lincoln, Nov 2017

U. Colorado, Boulder, Oct 2017

SMBE structural variation symposium, San Antonio, July 2017

Harris Moran Breeding, Davis, Feb 2017

PAG next-generation sequencing symposium, Jan 2017 (declined)

VI Brazilan Plant Molecular Genetics symposium, Ouro Preto, Brazil, May 2017 (declined)

Davis Science Cafe, Aug 2016

SMBE domestication symposium, Queensland, Australia, July 2016

UC Master Gardeners, May 2016

U. Arizona, Apr. 2016

Jo 6 conferencesGenome Institute, Mar. 2016

U. Southern California, Feb. 2016

LANGEBIO, Irapuato, Mexico, Nov. 2015

U. Toronto, Oct. 2015

Danforth Center, Sept. 2015

SMBE workshop on adaptation and next-gen sequencing, Montpellier, June 2015

San Francisco Exploratorium, May 2015

Dept. of Ecology and Evol. Bio, UC Irvine, April 2015

Cornell Plant Breeding Symposium, March 2015

LANGEBIO (Irapuato), Sept. 2014

Pioneer Hi-Bred (IA), Sept. 2014

Dept. of Ecology and Evolution, Iowa State U., Sept. 2014

Pioneer Hi-Bred (CA), Aug. 2014

Bioagricultural Sciences and Pest Management, Colorado State U., May 2014

Plant Breeding Genetics and Biotechnology Program, Michigan State (MI), Apr. 2014

National Maize Improvement Center of China, China Agricultural University (Beijing), Mar. 2014 Dept. of Agronomy, University of Guelph, Feb. 2014

Plant and Animal Genome Conference, maize workshop, Jan. 2014

Plant and Animal Genome Conference, symposium on domestication, Jan. 2014

Featured Speaker, Ecological Genomics Symposium, Ecological Genetics Institute (MO), Nov. 2013

Department of Genetics, U. Georgia, Sept. 2013

Plenary Speaker, Canadian Plant Genomics Workshop (Halifax) Aug. 2013

Organizer, Evolutionary Genomics symposium, ASPB (RI) 2013

Biodesign Institute, Arizona State U. 2013

Interdisciplinary Plant Group, U. Missouri 2013

UCD@BGI featured speaker, UC Davis 2013

Plant and Animal Genome Conference, symposium on translational genomics (CA) 2013

Featured Speaker, UC Davis Seed Central 2013

Crop Wild Relative Genomics meeting (CA) 2012

Germplasm Enhancement of Maize, ASTA Conference (IA) 2012

Pioneer Hi-Bred (CA) 2012

Plenary Speaker, Coastwide Salmonid Genomics Conference (CA) 2012

BASF Plant Science (NC) 2012

Pioneer Hi-Bred (IA) 2012

Illinois Corn Breeders School (IL) 2012

Keynote Speaker, North Central Regional Corn Breeding Research Meeting (IL) 2012

Plant and Animal Genome Conference, symposium on ecological genomics (CA) 2012

ASA/CSSA/SSSA Convention, symposium on maize biology (TX) 2011

Dept. of Plant & Microbial Biology, UC Berkeley 2011

Seminis Vegetable Seeds (CA) 2011

Dept. of Plant Sciences, UC Davis 2011

Center for Population Biology, UC Davis 2011

Dept. of Botany and Plant Sciences, UC Riverside 2011

USDA Agricultural Research Service, Iowa State U. 2010

Microbial and Plant Genomics Institute, U. Minnesota 2010

Society for Molecular Biology and Evolution, Plant Ecological Genomics Symposium (France) 2010

Dept. of Plant Sciences, UC Davis 2009

Instituto de Ecología, Universidad Nacional Autónoma de México 2008

Harlan II Symposium, UC Davis 2008

Dept. of Biology, UC Riverside 2008

Secretaría de Medio Ambiente y Recursos Naturales, GMO Risk Assessment (Mexico) 2008

Dept. of Plant Sciences, UC Davis 2007

Dept. of Biology, York University 2007

Dept. of Botany and Plant Sciences, UC Riverside 2007

Georgia Partnership for Reform in Science and Mathematics (PRISM), U. Georgia 2004

University of Georgia Chapter of Sigma-Xi, U Georgia 2004

Publications (lab members bold, *equal contribution, ‡undergraduate, §corresponding, [citations])

Preprints

Lozano R, Gazave E, dos Santos JPR, Stetter MG, Valluru R, Vandliio N, Fernandes SB, Brown PJ, Shakoor N, Mockler T, Ross-Ibarra J, Buckler ES, Gore M. Comparative evolutionary analysis and

prediction of deleterious mutation patterns between sorghum and maize. doi: 10.1101/777623 [1]

Gates DJ[§], Runcie D, Janzen GM, Romero Navarro A, Willcox M, Sonder K, Snodgrass SJ, Rodríguez-Zapata F, Sawers RJH, Rubén Rellín-Álvarez, Buckler ES, Hearne S, Hufford MB, **Ross-Ibarra J**[§]. Single-gene resolution of locally adaptive genetic variation in Mexican maize. doi: 10.1101/706739 [o]

Stitzer MC[§], Anderson SN, Springer NM, **Ross-Ibarra J**. The Genomic Ecosystem of Transposable Elements in Maize. doi: 10.1101/559922 [8]

In press or in print

H-Index 42 (8712 citations as of Fri May 8 14:43:31 2020)

- 91. Chen Q, Samayo LF, Yang CJ, Bradbury PJ, Olukolu BA, Neumeyer MA, Romay, MC, Sun Q, Lorant A, Buckler ES, Ross-Ibarra J, Holland JB, Doebley JF (2020). The genetic architecture of the maize progenitor, teosinte, and how it was altered during maize domestication PLoS GENETICS *Accepted* [o]
- 90. **Zeitler L**, **Ross-Ibarra J**§, **Stetter MGS**§ (2020). Loss of diversity and accumulation of genetic load in doubled-haploid lines from European maize landraces. G₃ *Accepted* [o]
- 89. Wang B, Lin Z, Li X, Zhao Y, Zhao B, Wu G, Ma X, Wang H, Xie Y, Li Q, Song G, Kong D, Zheng Z, Wei H, Shen R, Chen C, Meng Z, Wang T, Li X, Chen Y, Lai J, Hufford MB, Ross-Ibarra J, He H, Wang H (2020). Genome-wide selection and genetic improvement during modern maize breeding. Nature Genetics *In Press* [o]
- 88. Torres R*, **Stetter MG***, Hernandez R[§], **Ross-Ibarra J**[§] (2020). The temporal dynamics of background selection in non-equilibrium populations. Genetics 214: 1019-1030 [6]
- 87. **Turner-Hissong SD**[§], Mabrey ME, Beissinger TM, **Ross-Ibarra J**, Pires JC (2020). Evolutionary insights into plant breeding. Current Opinion in Plant Biology 54: 93-100 [0]
- 86. Anderson SN, **Stitzer MC**, Zhou P, **Ross-Ibarra J**, Hirsch CD, Springer NM (2019) Dynamic patterns of transcript abundance of transposable element families in maize. G₃ 9: 3673-3682 [4]
- 85. Anderson SN*, **Stitzer MC***, Brohammer A*, Zhou P, Noshay JM, O'Connor CH, Hirsch CD, **Ross-Ibarra J**, Hirsch CN, Springer NM (2019). Transposable elements contribute to dynamic genome content in maize. The Plant Journal 100: 1052-1065 [13]
- 84. Wei X, Anderson SN, Wang X, Yang L, Crisp PA, Li Q, Noshay J, Albert PS, Birchler JA, **Bilinski MC**, **Stitzer MC**, **Ross-Ibarra J**, Flint-Garcia S, Chen X, Springer NM, Doebley JF (2019). Hybrid decay: a transgenerational epigenetic decline in vigor and viability triggered in backcross populations of teosinte with maize. Genetics 213: 143-160 [0]
- 83. **O'Brien AM**[§], Sawers RJH, Strauss SY, **Ross-Ibarra J**[§] (2019). Adaptive phenotypic divergence in teosinte differs across biotic contexts. Evolution 73: 2230-2246 [o]
- 82. Gonzalez-Segovia E, Pérez-Limon S, Cíntora-Martínez C, Guerrero-Zavala A, Jansen G, Hufford MB, Ross-Ibarra J, Sawers RJH (2019). Characterization of introgression from the teosinte *Zea*

- mays ssp. mexicana to Mexican highland maize. Peer 7: e6815. [1]
- 81. **Josephs EM**§, Berg JJ, **Ross-Ibarra J**, Coop G (2019) Detecting adaptive differentiation in structured populations with genomic data and common gardens. Genetics 211: 989-1004. [12]
- 80. **Stetter MG**[§], Thornton K, **Ross-Ibarra J**[§] (2018) Genetic architecture and selective sweeps after polygenic adaptation to distant trait optima. PLoS GENETICS 14(11): e1007794. [13]
- 79. **O'Brien A** \S , Sawers R, **Ross-Ibarra J**, Strauss SY \S (2018) Evolutionary responses to conditionality in species interactions across environmental gradients. American Naturalist 192(6): 715-730. [o]
- 78. **Stitzer MC**[§], **Ross-Ibarra J** (2018) Maize domestication and gene interaction. New Phytologist 220:395-408 [22]
- 77. Manchanda N, Snodgrass SJ, **Ross-Ibarra J**, Hufford MB (2018) Evolution and adaptation in the maize genome. *In* The Zea Mays Genome, Bennetzen, Flint-Garcia, Hirsch, Tuberosa (Eds.), Springer Nature Publishing *In Press* [0]
- 76. **Lorant A**, **Ross-Ibarra J**, Maud Tenaillon (2018) Genomics of long- and short- term adaptation in maize and teosinte. *In* Statistical Population Genomics, Dutheil (Ed.), Springer Nature Publishing *In Press* [0]
- 75. Dawe RK, Lowry EG, Gent J, **Stitzer MC**, Higgins DM, **Ross-Ibarra J**, Wallace JG, Kanizay L, Alabady M, Wang N, Gao Z, Birchler J, Harkess AE, Hodges AL, Hiatt EN (2018) A novel maize kinesin causes neocentromere activity and meiotic drive, altering inheritance patterns across the genome. Cell 173: 839-850. [31]
- 74. Aburto-Oropeza O, Johnson A, Agha M, Allen E, Allen M, González JA, Arenas-Moreno DM, Beas R, Butterfield H, Caetano G, Caselle J, Casteñada Gaytán G, Castorani MCN, Anh Cat L, Cavanaugh K, Chambers JQ, Cooper RD, Arafeh-Dalmau N, Dawson T, Diaz de la Vega A, DiMento JFC, Domínguez S, Edwards M, Ennen J, Estrada-Medina H, Fierro N, Gadsden H, Galina-Tessaro P, Gibbons P, Goode EV, Gorris ME, Harmon T, Hecht SB, Heredia Fragoso MA, Hernández-Solano A, Hernández-Cortés D, Hernández-Carmona G, Hillard S, Huey RB, Hufford MB, Pàramo Figueroa VH, Jenerette D, Jiménez-Osornio J, López-Nava KJ, Lara R, Leslie H, Lopez-Feldman A, Luja V, Martínez-Méndez N, Mautz W, Medellin-Azuara J, Meléndez-Torres C, de la Cruz FRM, Micheli F, Miles D, Montagner G, Montaño-Moctezuma G, Müller J, Oliva P, Ortinez A, Ortiz Partida JP, Palleiro-Nayar J, Parnell PE, Raimondi P, Ramirez A, Randerson JT, Reed DC, Riquelme M, Torres TR, Rosen PC, Ross-Ibarra J, Sanchez-Cordero V, Sandoval-Solis S, Santos J, Sawers R, Sinervo B, Sites J, Sosa-Nishizaki O, Stanton T, Stapp J, Stewart J, Torre J, Torres-Moye G, Treseder KK, Valdez-Villavicencio JH, Jiménez FIV, Vaughn M, Welton L, Westphal MF, Woolrich-Piña G, Yunez-Naude A, Zertuche-González JA, Taylor JE (2018) Harnessing Cross-border Resources to Confront Climate Change. Environmental Science and Policy *In Press*. [3]
- 73. **Bilinski P**[§], Albert P, Berg JJ, Birchler JA, Grote M, **Lorant A**, **Quezada** J[‡], Swarts, K, **Yang J**, **Ross-Ibarra** J[§] (2018) Parallel altitudinal clines reveal adaptive evolution of genome size in *Zea mays*. PLoS Genetics 14: e1007162 [0]
- 72. **Mei W, Stetter MG, Gates DJ, Stitzer MC, Ross-Ibarra J**§ (2018) Adaptation in plant genomes: bigger is different. American Journal of Botany 105: 16-19 [26]
- 71. Bukowski R, Guo X, Lu Y, Zou C, He B, Rong Z, Wang B, Xu D, Yang B, Xie C, Fan L, Gao S, Xu X, Zhang G, Li Y, Jiao Y, Doebley J, Ross-Ibarra J, Lorant A, Buffalo V, Romay MC, Buckler ES,

- Ware D, Lai J, Sun Q, Xu Y (2017) Construction of the third generation *Zea mays* haplotype map. GigaScience gix134 [0]
- 70. Wang L, **Beissinger TM**, **Lorant A**, **Ross-Ibarra C**, **Ross-Ibarra J**[§], Hufford MB[§] (2017) The interplay of demography and selection during maize domestication and diffusion. Genome Biology 18:215 [62]
- 69. Yang J*§, Mezmouk S*, Baumgarten A, Buckler ES, Guill KE, McMullen MD, Mumm RH, Ross-Ibarra J[§] (2017) Incomplete dominance of deleterious alleles contribute substantially to trait variation and heterosis in maize. PLoS Genetics 13:e1007019 [50]
- 68. **Lorant A**, Pedersen S, Holst I, Hufford MB, Winter K, Piperno D, **Ross-Ibarra J**§ (2017) The potential role of genetic assimilation during maize domestication. PLoS ONE 12:e0184202 [11]
- 67. Aguilar-Rangel MR, Chàvez Montes RA, Gonzalez-Segovia E, **Ross-Ibarra J**, Simpson JK, Sawers RJH (2017) Allele specific expression analysis identifies regulatory variation associated with stress-related genes in the Mexican highland maize landrace Palomero Toluqueño. PeerJ 5:e3737 [10]
- 66. **Stetter MG**[§], **Gates DJ**, **Mei W**, **Ross-Ibarra J**[§] (2017) How to make a domesticate. Current Biology 27:R896-R900 [17]
- 65. Swarts K, Gutaker RM, Schuenemann V, Benz B, Blake M, Bukowski R, Holland J, Kruse-Peeples M, Lepak N, Matson RG, Prim L, Romay C, Ross-Ibarra J, Sanchez J, Schmidt C, Sofro E, Krause J, Weigel D, Buckler ES, Burbano HA (2017) Genomic estimation of complex traits reveals ancient maize adaptation to temperate North America. Science 357:512-515 [63]
- 64. **Bilinski P**[§], Han Y, **Hufford MB**, **Lorant A**, Zhang P, Jiang J, **Ross-Ibarra J**[§] (2017) Genomic abundance is not predictive of tandem repeat localization in grass genomes. PLoS ONE 12:e0177896 [6]
- 63. Jiao Y, Peluso P, Shi J, Liang T, **Stitzer MC**, Wang B, Campbell M, Stein JC, Wei X, Chin C-S, Guill K, Regulski M, Kumari S, Olson A, Gent J, Schneider KL, Wolfgruber TK, May MR, Springer N, Antoniou E, McCombie R, Presting GG, McMullen M, **Ross-Ibarra J**, Dawe RK, Hastie A, Rank DR, Ware D (2017) Improved maize reference genome with single-molecule technologies. Nature 546:524-527 [397]
- 62. **Renny-Byfield S**§, Rodgers-Melnick E, **Ross-Ibarra J**§ (2017) Gene fractionation and function in the ancient subgenomes of maize. MBE 34:1825-1832 [29]
- 61. **Velasco D**, Aradhya M, and and **Ross-Ibarra J**§ (2016) Evolutionary genomics of peach and almond domestication. G₃ 6:3985-3993 [22]
- 60. Ramos-Madrigal J, Smith BD, Moreno-Mayar JV, Gopalakrishnan S, **Ross-Ibarra J**, Gilbert MTP, Wales N (2016) Genome sequence of a 5310-year-old maize cob provides insights into the early stages of maize domestication. Current Biology 26:3195-3201 [22]
- 59. **Durvasula A**^{‡*}, Hoffman PJ*, **Kent TV**[‡], Liu C, Kono TJY, Morrell PL[§], **Ross-Ibarra J**[§] (2016) ANGSD-wrapper. Molecular Ecology Resources 16:1449-1454 [o]
- 58. **Beissinger TM**§, Wang L, **Crosby K**, **Durvasula A**‡, Hufford MB, **Ross-Ibarra J**§ (2016) Recent demography drives changes in linked selection across the maize genome. Nature Plants 2:16084 [75]

57. Wolfgruber TK, Nakashima MM, Schneider KL, Sharma A, Xie Z, Albert PS, Xu R, **Bilinski P**, Dawe RK, **Ross-Ibarra J**, Birchler JA, Presting G (2016) High quality maize centromere 10 sequence reveals evidence of frequent recombination events. Frontiers In Plant Science 7 [22]

- 56. Orozco-Ramìrez Q, Santacruz-Varela A, **Ross-Ibarra J**, Brush B (2016) Maize diversity associated with social origin and environmental variation in southern Mexico. Heredity 116:477-484. [31]
- 55. Gerke JP[§], Edwards JW, Guill KE, **Ross-Ibarra J**[§], McMullen MD. The genomic impacts of drift and selection for hybrid performance in maize (2015). GENETICS 201: 1201-1211 [30]
- 54. Sosso D, Luo D, Li Q-B, Sassse J, Yang J, Gendrot G, Suzuki M, Koch KE, McCarty DR, Chourey PS, Rogoswky PM, Ross-Ibarra J, Yang B, Frommer WB (2015) Seed filling in domesticated maize and rice depends on SWEET-mediated hexose transport. Nature Genetics 47:1489-1493 [130]
- 53. **Takuno S**, Ralph P, Swarts K, Elshire RJ, Glaubitz JC, Buckler ES, **Hufford MB**, **Ross-Ibarra** J[§] (2015) Independent molecular basis of convergent highland adaptation in maize. Genetics 200:1297-1312 [50]
- 52. **Vann LE**, **Kono T**, **Pyhäjärvi T**, **Hufford MB**[§], **Ross-Ibarra J**[§] (2015) Natural variation in teosinte at the domestication locus teosinte branched1 (tb1). PEERJ 3:e900 [11]
- 51. Hake S, **Ross-Ibarra J** (2015) Genetic, evolutionary and plant breeding insights from the domestication of maize. ELIFE 2015;4:e05861 [39]
- 50. Fonseca RR, Smith B, Wales N, Cappellini E, Skoglund P, Fumagalli M, Samaniego JA, Caroe C, Avila-Arcos MC, Hufnagel D, Korneliussen TS, Vieira FG, Jakobsson M, Arriaza B, Willerslev E, Nielsen R, Hufford MB, Albrechtsen A, Ross-Ibarra J, Gilbert MT (2015) The origin and evolution of maize in the American Southwest. Nature Plants 1:14003 [100]
- 49. Dyer GA, López-Feldman A, Yúnez-Naude A, Taylor JE, **Ross-Ibarra J** (2015) Reply to Brush *et al.*: A wake up call for crop conservation science. PNAS 112 (1), E2-E2 (letter). [7]
- 48. Makarevitch I, Waters M, West P, **Stitzer M**, **Ross-Ibarra**, **J**, Springer NM (2015) Mobile elements contribute to activation of genes in response to abiotic stress. PLoS Genetics 11 (1): e1004915. [223]
- 47. Tiffin P, **Ross-Ibarra J** (2014) Advances and limits of using population genetics to understand local adaptation. Trends in Ecology and Evolution 29:673-680 [223]
- 46. **Bilinski P, Distor KD, Gutierez-Lopez J, Mendoza Mendoza G**, Shi J, Dawe RK, **Ross-Ibarra J**[§] (2014) Diversity and evolution of centromere repeats in the maize genome. Сняомозома 0009-5915 [17]
- 45. **Mezmouk S**, **Ross-Ibarra J** § (2014) The pattern and distribution of deleterious mutations in maize. (2014). G₃ 4:163-171 [61]
- 44. Waters AJ, **Bilinski P**, Eichten SR, Vaughn MW, **Ross-Ibarra J**, Gehring M, Springer NM (2013) Comprehensive analysis of imprinted genes in maize reveals allelic variation for imprinting and limited conservation with other species. PNAS 110:19639-19644 [83]
- 43. **Pyhäjärvi T**, **Hufford MB**, **Mezmouk S**, **Ross-Ibarra J**[§] (2013) Complex patterns of local adaptation in teosinte. Genome Biology and Evolution 5: 1594-1609 [106]

42. Wills DM, Whipple C, **Takuno S**, Kursel LE, Shannon LM, **Ross-Ibarra J**, Doebley JF (2013) From many, one: genetic control of prolificacy during maize domestication. PLoS GENETICS 9(6): e1003604. [74]

- 41. McCouch S, Baute GJ, Bradeen J, Bramel P, Bretting PK, Buckler E, Burke JM, Charest D, Cloutier S, Cole G, Dempewolf H, Dingkuhn M, Feuillet C, Gepts, P, Grattapaglia D, Guarino L, Jackson S, Knapp S, Langridge P, Lawton-Rauh A, Lijua Q, Lusty C, Michael T, Myles S, Naito K, Nelson RL, Pontarollo R, Richards CM, Rieseberg L, Ross-Ibarra J, Rounsley S, Hamilton RS, Schurr U, Stein N, Tomooka N, van der Knaap E, van Tassel D, Toll J, Valls J, Varshney RK, Ward J, Waugh R, Wenzl P, Zamir. (2013) Agriculture: Feeding the future. Nature 499:23-24 [334]
- 40. **Hufford MB**, Lubinsky P, **Pyhäjärvi T**, **Devengenzo MT**[‡], Ellstrand NC, **Ross-Ibarra J**[§] (2013) The genomic signature of crop-wild introgression in maize. PLoS Genetics 9(5): e1003477. [219]
- 39. **Provance MC**[§], Garcia Ruiz I, **Thommes C**[‡], **Ross-Ibarra J** (2013) Population genetics and ethnobotany of cultivated *Diospyros riojae* Gómez Pompa (Ebenaceae), an endangered fruit crop from Mexico. Genetic Resources and Crop Evolution 60: 2171-2182. [5]
- 38. Melters DP*, Bradnam KR*, Young HA, Telis N, May MR, Ruby JG, Sebra R, Peluso P, Eid J, Rank D, Fernando Garcia J, DeRisi J, Smith T, Tobias C, **Ross-Ibarra J**[§], Korf IF[§], Chan SW-L. (2013) Patterns of centromere tandem repeat evolution in 282 animal and plant genomes. Genome Biology 14:R10 [226]
- 37. Kanizay LB, **Pyhäjärvi T**, Lowry E, **Hufford MB**, Peterson DG, **Ross-Ibarra J**, Dawe RK (2013) Diversity and abundance of the Abnormal chromosome 10 meiotic drive complex in *Zea mays*. Heredity 110: 570-577. [19]
- 36. **Hufford MB**, **Bilinski P**, **Pyhäjärvi T**, **Ross-Ibarra J**§ (2012) Teosinte as a model system for population and ecological genomics. Trends in Genetics 12:606-615 [51]
- 35. Muñoz Diez C, Vitte C, **Ross-Ibarra J**, Gaut BS, Tenaillon MI (2012) Using nextgen sequencing to investigate genome size variation and transposable element content. *In* Grandbastien M-A, Casacuberta JM, editors. Topics in Current Genetics v24: Plant Transposable Elements Impact on Genome Structure & Function. pp. 41-58 [18]
- 34. **van Heerwaarden J**§, **Hufford MB**, **Ross-Ibarra J**§ (2012) Historical genomics of North American maize. PNAS 109: 12420-12425 [111]
- 33. Swanson-Wagner R, Briskine R, Schaefer R, **Hufford MB**, **Ross-Ibarra J**, Myers CL, Tiffin P, Springer NM. Reshaping of the maize transcriptome by domestication. (2012). PNAS 109: 11878-11883 [112]
- 32. **Hufford MB***, Xun X*, **van Heerwaarden J***, **Pyhäjärvi T***, Chia J-M, Cartwright RA, Elshire RJ, Glaubitz JC, Guill KE, Kaeppler S, Lai J, Morrell PL, Shannon LM, Song C, Spinger NM, Swanson-Wagner RA, Tiffin P, Wang J, Zhang G, Doebley J, McMullen MD, Ware D, Buckler ES[§], Yang S[§], **Ross-Ibarra J**[§] (2012) Comparative population genomics of maize domestication and improvement. Nature Genetics 44:808-811 [578]
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