## curriculum vitae DEREN A. R. EATON

Columbia University

Department of Ecology, Evolution, and Environmental Biology

Schermerhorn Extension 1007 New York, NY 10027, USA Email: de2356@columba.edu

**Appointments** 

7/2017 - Assistant Professor, Columbia University, Ecology, Evolution, &

**Environmental Biology** 

5/2016 - 7/2017 Associate Research Scientist, Yale University, Ecology and

**Evolution** 

1/2014 - 5/2016 Postdoctoral researcher, Yale University, Ecology and Evolution

Education

9/2008 - 12/2013 Ph.D. University of Chicago, Committee on Evolutionary Biology

9/2003 - 5/2007 B.S. University of Minnesota, Plant Biology (Honors)

9/2003 - 5/2007 B.S. University of Minnesota, Ecology, Evolution and Behavior

#### **Grants and Awards**

#### *Grants Awarded:*

Field course; Principles and Practice of Modern Genomic Data Science

Source: Earth Institute Course Support Grant for travel expenses

Duration: 1/2019 - 5/2019

Funding: \$4,000

Role: PI

Principles and Practice of Modern Genomic Data Science

Source: Columbia University Center for Teaching and Learning Hybrid Learning Grant

Duration: 9/2018 - 5/2019

Funding: \$13,400

Role: PI

Genomic and Chemical Consequences of Different Levels of Gene Flow in the Fig-Wasp

Pollination Mutualism

Source: Smithsonian Scholarly Studies Award

Duration: 9/2016 Funding: \$50,324 Role: Co-PI

Replicated Evolution of Leaf Form in a Neo-tropical Radiation of Viburnum

Source: NSF DEB 1557059

Funding: \$996,515

Duration: 5/2016 - 5/2019

Role: Co-PI

Comparative Studies in Reproductive Interference: Linking Floral Adaptations and Species

#### Divergence.

Source: NSF Doctoral Dissertation Improvement Grant

Funding: \$15,000

Duration: 5/2011 – 5/2014

Role: PI

## **Grants Contributed to:**

The Evolutionary Roles of Hybridization and Introgression: Investigating Species and

Genomic Boundaries Under Climate Change; Primula Genomics: PrimGEN

Source: Swiss National Science Foundation

Duration: 5/2018 – 5/2022 Funding: CHF 700,000

Role: Personnel

Dimensions: Tropical Niche Conservatism in Drosophila: Testing the Genetic and

Functional Constraints on Diversification

Source: NSF DEB 1737752 Duration: 5/2017 - 5/2020 Funding: \$1,946,771 Role: Personnel

### Fellowships:

9/2011 - 9/2012 Lester Armour Graduate Fellowship, Field Museum; \$30,000 6/2010 - 9/2010 NSF East Asian Pacific Summer Institutes, China; \$5,600

#### **Small Awards:**

| 2015      | Yale Postdoctoral scholars travel award                                 |
|-----------|---|
| 2012      | Micromorph workshop, Harvard, travel and board                          |
| 2011      | Hinds Fund Grant, University of Chicago                                 |
| 2009      | Pritzker Laboratory Grant, The Field Museum                             |
| 2008      | NSF Graduate research fellowship; Honorable Mention                     |
| 2007      | Honors, summa cum laude, U Minnesota                                    |
| 2007      | Ernst Abbe Award for Plant Biology Majors, U Minnesota                  |
| 2005-2007 | Dean's List scholarship, U Minnesota                                    |
| 2006      | Center for International Education Exchange (CIEE) Student of the Year  |
| 2004-2007 | Soil, Water, Climate Department scholarship, U Minnesota                |
| 2004-2006 | College of Agriculture, Food and Environmental Science scholarship, UMN |
|           |   |

#### **Publications**

2019 Satler, J.D., Herre, A., Jander, C., **Eaton, D.A.R.,** Machado, C.A., Heath, T.A., and J.D. Nason (2019). *Inferring Processes of Coevolutionary Diversification in a Community of Panamanian Strangler Figs and Associated Pollinating Wasps*. BioRxiv, December, 490862. https://doi.org/10.1101/490862.

2019 Spriggs, E.L., **Eaton, D.A.R.,** Sweeney, P., Schlutius, C., Edwards, E., and M.J. Donoghue. (2019) *RAD-seq data reveal a cryptic Viburnum species on the North* 

- American Coastal Plain. American Journal of Botany. In Press.
- Miller, J., Quinzin, M., Edwards, **D., Eaton, D.A.R.**, Jensen, E., Russello, M., Gibbs, J., Tapia, W., Rueda, D., and A. Caccone. (2018). *Genome-wide assessment of diversity and divergence among extant Galápagos giant tortoise species*. Journal of Heredity. doi: 10.1093/jhered/esy031.
- 2018 Park, B., Sinnott-Armstrong, M., Schlutius, C., Penagos Zuluaga, J., Spriggs, E.L, Simpson, R., Benavides, E., Landis, M., Sweeney, P., **Eaton, D.A.R.**, and M.J. Donoghue. (2018). *Sterile marginal flowers increase visitation and fruit set in the hobblebush (Viburnum lantanoides, Adoxaceae) at multiple spatial scales*. Annals of Botany; doi: 10.1093/aob/mcy117
- 2018 Federman, S., Donoghue, M.J., Daly, D., and **D.A.R. Eaton.** (2018). *Reconciling Species Diversity in a Tropical Plant Clade (Canarium, Burseraceae).* PLoS One 13(6): e0198882.
- 2018 McKain, M.R., Johnson, M.G., Uribe-Convers, S., **Eaton, D.A.R.,** and Y. Yang \*(*All authors contributed equally*) (2018). *Practical Considerations for Plant Phylogenomics*. Applications in Plant Sciences 6(3):e1038.
- 2017 Forsman, Z.H., Knapp, I.S.S, Tisthammer, K., Eaton, D.A.R., Belcaid, M., and R.J Toonen (2017). Coral hybridization or phenotypic variation? Genomic data reveal gene flow between Porites lobata and P. compressa. Molecular Phylogenetics and Evolution 111:132-148.
- 2016 **Eaton, D.A.R.,** Spriggs, E.L., Park, B. and M.J. Donoghue (2016). *Misconceptions on missing data in RADseq phylogenetics with a deep-scale example from flowering plants (Viburnum: Adoxaceae*). Systematic Biology (https://doi.org/10.1093/sysbio/syw092).
- 2015 **Eaton, D.A.R.,** Hipp, A., Gonzalez-Rodriguez, A. and J. Cavender-Bares (2015). *Historical introgression among the American live oaks and the comparative nature of tests for introgression.* Evolution 69(10): 2587-2601.
- 2015 Cavender-Bares, J., Gonzalez-Rodriguez, A., **Eaton, D.A.R.**, Hipp, A., Buelke, A., and P. Manos (2015). *Phylogeny and biogeography of the American live oaks* (*Quercus subsection Virentes*): *A genomic and population genetic approach*. Molecular Ecology, 24(14): 3668-3687
- 2014 Escudero, M., **Eaton, D.A.R.**, Hahn, M. and A. Hipp (2014). *Genotyping-by-sequencing as a tool for phylogenetic inference and testing ancestral hybridization: A case study in Carex (Cyperaceae*). Molecular Phylogenetics and Evolution 79: 359-367.
- 2014 **Eaton, D.A.R.** (2014). *On the Evolutionary Consequences of Interspecific Reproductive Interactions.* Ph.D. Dissertation. University of Chicago.

- **Eaton, D.A.R.** (2014). *PyRAD: de novo Assembly of RAD/GBS data for phylogenetic and introgression analyses.* Bioinformatics, 30(13): 1844-1849.
- 2014 Hipp, A., **Eaton, D.A.R.**, Cavender-Bares, J., Fitzek, E., Nipper, R. and P. Manos (2014). *A framework phylogeny of the New World oak clade based on sequenced RAD data*. PLoS ONE 9(4): e93975.
- 2013 **Eaton, D.A.R.** and R.H. Ree (2013). *Inferring Phylogeny and Introgression using genomic RADseq Data: An Example from Flowering Plants (Pedicularis: Orobanchaceae)*. Systematic Biology, 62: 689-706
- Wang, X., Zhao, L., **Eaton, D.A.R.** and Z. Guo (2013). *Identification of SNP markers for inferring phylogeny in temperate Bamboos (Poaceae: Bambusoideae) using RAD tag sequencing*. Molecular Ecology Resources, 13: 938-945.
- 2013 Fournier-Level A., Wilczek, A.M., Cooper, M.D., Roe, J.L, Anderson, J.A., **Eaton, D.A.R.**, Moyers, B.T., Petipas, R.H., Schaeffer, R.N., Pieper, B., Reymond, M., Koorneef, M., Welch, S.M., Remington, D.L. and J.S. Schmitt (2013). *Paths to selection on life-history loci in different natural environments across the native range of Arabidopsis thaliana*. Molecular Ecology, 22: 3552-3566.
- 2012 **Eaton, D.A.R.**, Fenster, C.B., Hereford, J., Huang, S-Q. and R.H. Ree (2012). *Floral diversity and community structure in Pedicularis (Orobanchaceae)*. Ecology, 93: S182-S19

### Other publications (non peer-reviewed)

2013 Hipp, A.L., Manos, P.S., Cavender-bares, J.C., **Eaton, D.A.R.**, and R. Nipper *Using Phylogenomics to infer the evolutionary history of Oaks*. The Journal of International Oaks. International Oak Journal, 24: 61-71

#### **Presentations**

| <b>Invited talks:</b> |   |
|-----------------------|---|
| 3/2019                | CUNY City College – Biology Department Seminar                          |
| 2/2019                | University of Wisconsin Madison – J.F. Crow Evolution Institute Seminar |
| 2/2019                | University of Wisconsin Madison – Darwin Day Public Invited Speaker     |
| 11/2018               | CUNY City Tech – Bioinformatics Colloquium Seminar Series               |
| 11/2018               | Pace University – Biology Department Seminar                            |
| 7/2018                | Kunming Institute of Botany, Chinese Academy of Sciences                |
| 5/2018                | Universidad Nacional Autonoma de Mexico, Mexico City                    |
| 4/2018                | Columbia University – Evolutionary Genomics Supergroup                  |
| 10/2017               | Missouri Botanic Garden – Annual Fall Symposium                         |
| 8/2017                | Workshop on Molecular Evolution – Marine Biological Laboratory          |
| 4/2017                | University of Michigan – EEB Department Seminar                         |
| 4/2016                | Universidad Nacional Autonoma de Mexico, Morelia – Invited seminar      |
| 4/2016                | NSF-NSFC China-US biodiversity workshop, Zhejiang University, China     |
| 3/2016                | University of Arkansas – Graduate Student Elected Invited Speaker       |
| 3/2016                | Rancho Santa Ana Botanic Garden – Graduate Student Elected Speaker      |
| 2/2016                | University of Idaho – IBEST RAD-seq symposium                           |

| 2/2016  | University of Minnesota – Plant Biology Department Seminar              |
|---------|---|
| 2/2016  | Columbia University – E3B Department Seminar                            |
| 10/2015 | Amherst College – Biology Department Seminar                            |
| 10/2015 | American Museum of Natural History – Comparative Biology Seminar        |
| 7/2015  | Universidad Nacional Autonoma de Mexico, D.F.                           |
| 3/2015  | Harvard University – C. Davis Lab seminar                               |
| 3/2015  | University of Hawaii – Department seminar                               |
| 3/2015  | Hawaii Institute of Marine Biology – Department seminar                 |
| 2/2014  | Yale University – EEB Department Seminar                                |
| 6/2013  | Ernst Mayr Symposium – Evolution Meeting, Snowbird, Utah                |
| 4/2013  | Yale University – Donoghue/Near lab seminar                             |
| 1/2013  | Field Museum – Chicago Plant Science Symposium                          |
| 12/2012 | University of Illinois at Chicago, Ecology and Evolution Dept. Seminar  |
| 8/2012  | Association for Tropical Biology and Conservation Conf., Bonito, Brazil |
| 8/2012  | Arnold Arboretum of Harvard University, microMORPH workshop             |
| 7/2010  | Shangri-la Alpine Botanic Garden, Yunnan, China, S-Q. Huang Lab         |

## Talks at Meetings/Conferences

| 1/2019  | International Biogeography Conference, Malaga, Spain                       |
|---------|--|
| 7/2016  | Botany meeting, Savannah, Georgia  |
| 6/2016  | Evolution meeting, Austin, Texas   |
| 7/2015  | Botany meeting, Edmonton, Alberta  |
| 1/2015  | Society for Systematic Biologists stand-alone meeting, Ann Arbor, Michigan |
| 7/2014  | Botany meeting, Boise, Idaho   |
| 6/2014  | Evolution meeting, Raleigh, North Carolina                                 |
| 6/2013  | Evolution meeting, Snowbird, Utah  |
| 6/2012  | Evolution meeting, Ottawa, Canada  |
| 11/2012 | Field Museum Plant Sciences Symposium                                      |
| 6/2012  | Field Museum Pritzker Laboratory Seminar                                   |
| 6/2009  | Evolution meeting, Moscow, Idaho   |

## Teaching (courses)

| 2019 Spring | Instructor, GR6300 Research Seminar, Columbia University                  |
|-------------|---|
| 2019 Spring | Instructor, GR4055 Principles and Applications of Modern DNA              |
|             | Sequencing, Columbia University   |
| 2018 Fall   | Instructor, GR6110 Fundamentals of Evolution, Columbia University         |
| 2018 Fall   | Instructor, GR6300 Research Seminar, Columbia University                  |
| 2018 Spring | Instructor, GR4050 Programming and Data Science for Biology, Columbia     |
|             | University  |
| 2017 Fall   | Instructor, GR6110 Fundamentals of Evolution, Columbia University         |
| 2013 Spring | Teaching Assistant, Ecology, Evolution of the Southwest Deserts, UChicago |
| 2011 Spring | Teaching Assistant, Environmental Ecology, UChicago                       |
| 2009 Spring | Teaching Assistant, Ecology, Evolution of the Southwest Deserts, UChicago |
| 2009 Summer | Teaching Assistant, Field Course in Desert Ecology, U of Chicago          |
| 2007 Spring | Teaching Assistant, Introduction to Biochemistry, U of Minnesota          |
|             |   |

# Organized Workshops: "Instructor: reproducible genomics workshop"

1/2019 International Biogeography Conference, Malaga, Spain

| 10/2018 | RADCamp 3-day genomics workshop, Columbia University, New York |
|---------|--|
| 11/2017 | Universidad Nacional Autonoma de Mexico, D.F.                  |
| 8/2017  | Workshop on Molecular Evolution, Woods Hole, MA                |
| 4/2017  | University of Oldenburg, Germany                               |
| 4/2016  | Universidad Nacional Autonoma de Mexico, Morelia               |
| 7/2016  | Botany conference, Savannah, Georgia                           |
| 3/2016  | University of Arkansas   |
| 3/2016  | Rancho Santa Ana Botanic Garden                                |
| 2/2016  | University of Idaho  |
| 5/2015  | Universidad Nacional Autonoma de Mexico, D.F.                  |

#### **Professional Societies**

Society of Systematic Biologists (SSB) Society for the study of Evolution (SSE) Botanical Society of America (BSA)

## Scientific Software (github.com/dereneaton, github.com/eaton-lab)

simcat – machine learning analysis of multidimensional phylogenetic invariants

strange – species tree and gene tree analysis in sliding windows

toytree – interactive tree plotting library for Python

*tetrad* – quartet based super tree inference using phylogenetic invariants

*ipyrad* – interactive assembly and analysis of RAD-seq data sets

*pyRAD* – assembly of RAD and GBS data sets

simrrls – simulate RADseq reads for analysis of mutation-disruption

*simloci* – simulate RADseq loci for analysis of mutation-disruption

## Signature

Our Exton

4/22/2019