

## Michael James Landis

[michael.landis@wustl.edu](mailto:michael.landis@wustl.edu)

[landislab.org](http://landislab.org)

Last updated: Nov 17, 2022

### Education

- 2015 Ph.D. Integrative Biology  
*Designated Emphasis in Computational & Genomic Biology*  
University of California, Berkeley, CA
- 2005 B.S. Computer Science  
California State University, Long Beach, CA

### Employment

- 2019— Assistant Professor  
Department of Biology  
Washington University, St. Louis, MO

### Research experience

- 2016—2019 Postdoctoral fellow, Yale University, New Haven, CT
- 2016 Postdoctoral researcher, Iowa State University, Ames, IA
- 2010—2015 Graduate researcher, University of California, Berkeley, CA
- 2009—2010 Bioinformatician, Children’s Hospital of Oakland, Oakland, CA

### Research Interests

I research relationships between evolutionary processes and patterns. My current interests include statistical phylogenetics, historical biogeography, the evolution of ecological interactions, phenotypic evolution, pathogen host switching, stochastic processes, model design, Bayesian inference, and deep learning. My work typically involves the design of probabilistic models and inference methods to test evolutionary hypotheses in biological and simulated datasets.

### Fellowships & Awards

- 2019 Yale Donnelley Postdoctoral Environmental Fellowship (resumed)
- 2017—2018 NSF Postdoctoral Research Fellowship
- 2016 Yale Donnelley Postdoctoral Environmental Fellowship
- 2016 Ernst Mayr Award (Society of Systematic Biologists)
- 2015 UC Berkeley Integrative Biology Summer Research Award
- 2012, 2015 UC Berkeley Computational Biology Travel Award
- 2013 National Evolutionary Synthesis Center Graduate Student Fellowship
- 2012 Google Summer of Code Student Fellowship

### External Grants

- 2021—2024 NSF-DEB 2040347: “Modeling the Origin and Evolution of Hawaiian Plants”  
Awarded \$547,408 to MJL as Lead PI
- 2022 WUSTL ITF: “The Human-Wildlife Interface: Disease Dynamics and Pandemic Prevention”  
Awarded \$116,000 to PIs Landis, Milich, Wang

### Manuscripts – Published

‡: group postdoc co-author; \*: shared first author

- 2022 1. Dismukes W, ‡Braga MP, Hembry DH, Heath TA, Landis MJ. Cophylogenetic methods to untangle the evolutionary history of ecological interactions. Accepted at Annual Reviews of Ecology, Evolution, and Systematics. 53:275–298.

2. Donoghue MJ, Eaton DAR, Maya-Lastra CA, **Landis MJ**, Sweeney PJ, Olson, ME, Cacho NI, Moeglin MK, Gardner JR, Heaphy NM, Castorena M, Segovia Rivas A, Clement WL, Edwards EJ. 2022. Replication radiation of a plant clade along a cloud forest archipelago. *Nature Ecology and Evolution*. doi:10.1038/s41559-022-01823-x.
3. Barido-Sottani J, Justison JA, Borges R, Brown JM, Dismukes W, do Rosario Petrucci B, Fabreti Guimarães L, Höhna S, **Landis MJ**, Lewis PO, May MR, Mendes FK, Pett W, Redelings BD, Tribble CM, Wright AM, Zenil-Ferguson R, Heath TA. 2022. Lessons learned from organizing and teaching virtual phylogenetics workshops. *Bulletin of the Society of Systematic Biologists* doi:10.18061/bssb.v1i2.
4. Wendt EW, Malabarba LR, **‡Braga MP**, Boeger WA, **Landis MJ**, Carvalho TP. Phylogeny, species delimitation, and ecological and morphological diversity of Characithecium (Monogeneoidea: Dactylogyridae). 2022. *Parasitology* 149: 700–716.
5. **Landis MJ**, Quintero I, Muñoz MM, Zapata F, Donoghue MJ. Phylogenetic inference of where species spread or split across barriers. 2022. *Proceedings of the National Academy of Sciences* 119: e2116948119.
6. Tribble CM, Freyman WA, Lim JY, **Landis MJ**, Barido-Sottani J, Kopperud BT, Höhna S, May MR. 2022. RevGadgets: an R Package for visualizing Bayesian phylogenetic analyses from RevBayes. *Methods in Ecology and Evolution* 13: 314–323.
- 2021 7. Höhna S, **Landis MJ**, and Huelsenbeck JP. 2021. Parallel power posterior analyses for fast computation of Bayes factors in phylogenetics. *PeerJ*: 9:e12438.
8. **‡Braga MP**, Janz N, Nylin S, Ronquist F, and **Landis MJ**. 2021. Phylogenetic reconstruction of ancestral ecological networks through time for pierid butterflies and their host plants. *Ecology Letters* 24: 2134–2145.
9. **Landis MJ**, Edwards EJ, and Donoghue MJ. 2021. Modeling phylogenetic biome shifts on a planet with a past. *Systematic Biology*. 70: 86–107.
10. **Landis MJ**, Eaton DAR, Clement WL, Park B, Spriggs EL, Sweeney PW, Edwards EJ, and Donoghue MJ. 2021. Joint phylogenetic estimation of geographic movements and biome shifts during the global diversification of Viburnum. *Systematic Biology*. 70: 67–85.
- 2020 11. **‡Braga MP**, **Landis MJ**, Nylin S, Janz N, and Ronquist F. 2020. Bayesian inference of ancestral host-parasite interactions under a phylogenetic model of host repertoire evolution. *Systematic Biology* 69: 1149–1162.  
*Awarded Best Paper in Systematic Biology in 2020.*
12. Field DJ, Berv JS, Hsiang AY, Lanfear J, **Landis MJ**, Dornberg A. 2020. Timing the extant avian radiation: The rise of modern birds, and the importance of modeling molecular rate variation. *Bulletin of the American Museum of Natural History* 440: 159–181.
13. Kim AS, Zimmerman O, Nelson CA, Basore K, Zhang R, Desai C, Bullock C, Durnell L, Deem SL, Oppenheimer J, Shapiro B, Wang T, Coyne CB, Handley SA, **§Landis MJ**, **§Fremont DH**, and **§Diamond MS**. 2020. A sequence insertion in the Mxra8 receptor of Bovinae family members confers resistance to alphavirus infection. *Cell Host & Microbe* 27: 428–440.
14. Quintero I and **Landis MJ**. 2020. Interdependent phenotypic and biogeographic evolution driven by biotic interactions. *Systematic Biology* 69: 739–755.
- 2018 15. **Landis MJ**, Freyman WA, and Baldwin BG. 2018. Retracing the Hawaiian silversword radiation despite phylogenetic, biogeographic, and paleogeographic uncertainty. *Evolution* 72: 2343–2359.
16. Park B, Sinnott-Armstrong M, Schlutius C, Zuluaga, J-CP, Spriggs EL, Simpson RG, **Landis MJ**, Sweeney PW, Eaton DAR, and Donoghue MJ. 2018. Sterile marginal flowers increase visitation and fruit set in the hobblebush (*Viburnum lantanoides*, Adoxaceae) at multiple spatial scales. *Ann. Bot.* 123: 381–390.

- 2017 17. **Landis MJ** and Schraiber JG. 2017. Pulsed evolution shaped modern vertebrate body sizes. *Proceedings of the National Academy of Sciences* 114: 13224–13229.
18. Höhna S, **Landis MJ**, Heath TA. 2017. Phylogenetic inference using RevBayes. *Current Protocols in Bioinformatics* 57:6.16.1–6.16.34.
19. **Landis, MJ**. Biogeographic dating of speciation times using paleogeographically informed processes. 2017. *Systematic Biology* 66:128–144.
- 2016 20. Höhna S, **Landis MJ**, Heath TA, Boussau B, Lartillot N, Moore BR, Huelsenbeck JP, and Ronquist F. 2016. RevBayes: Bayesian phylogenetic inference using graphical models and an interactive model-specification language. *Systematic Biology* 65:726–736.
- 2015 21. Schraiber JG and **Landis MJ**. 2015. Sensitivity of quantitative traits to mutational effects and number of loci. *Theoretical Population Biology* 102:85–93.
- 2014 22. Höhna S, Heath TA, Boussau B, **Landis MJ**, Ronquist F, and Huelsenbeck JP. 2014. Probabilistic graphical model representation in phylogenetics. *Systematic Biology* 63:753–771.
23. **Landis MJ** and Bedford T. 2014. Phylowood: interactive web-based animations of biogeographic and phylogeographic histories. *Bioinformatics* 30:123–124.
- 2013 24. **Landis MJ**, Matzke NJ, Moore BR, and Huelsenbeck JP. Bayesian analysis of biogeography when the number of areas is large. 2013. *Systematic Biology* 62:789–804.
25. \***Landis MJ**, \*Schraiber JG, and Liang M. 2013. Phylogenetic analysis using Lévy processes: finding jumps in the evolution of continuous traits. *Systematic Biology* 62:193–204.

#### Manuscripts – Under Review

‡: group postdoc co-author

- 2022 1. Kawahara AY, Storer C, Carvalho APS, Plotkin DM, Condamine F, ‡**Braga MP**, ..., **Landis MJ**, ..., Lohman DJ. Evolution and diversification dynamics of butterflies.
2. Nielsen SV, Vaughn A, Leppälä, **Landis MJ**, Mailund T, Nielsen R. Bayesian inference of admixture graphs on Native American and Arctic populations.

#### Book Chapters – Published

- 2020 1. Landis MJ. Biogeographic dating of phylogenetic divergence times using priors and processes. 2020. In Ho SYW (ed.), *The Molecular Evolutionary Clock: Theory and Practice*. Springer.

#### Research Software

RevBayes, phylogenetic inference using graphical models	<a href="https://revbayes.github.io">revbayes.github.io</a>
BayArea, Bayesian biogeographic inference for many areas	<a href="https://github.com/mlandis/bayarea">github.com/mlandis/bayarea</a>
pulsR, simulate and fit macroevolutionary trait models	<a href="https://github.com/Schraiber/pulsR">github.com/Schraiber/pulsR</a>
creepy-jerk, Bayesian inference of evolutionary jumps in traits	<a href="https://github.com/mlandis/creepy-jerk">github.com/mlandis/creepy-jerk</a>
Phylowood, interactive biogeographic animations	<a href="https://mlandis.github.io/phylowood">mlandis.github.io/phylowood</a>
qtc, trait evolution under the coalescent	<a href="https://github.com/Schraiber/quant_trait_coalescent">github.com/Schraiber/quant_trait_coalescent</a>

#### Presentations – Invited

\*: graduate student invited

- 2022 Meeting of Systematics, Biogeography, and Evolution (virtual)  
 \*Melinda Denton Endowed Seminar on Plant Systematics, University of Washington  
 \*Ecology and Evolution Seminar, University of Minnesota, Minneapolis (virtual)  
 Biodiversity Research Center Seminar, University of British Columbia, Vancouver (virtual)
- 2021 European Society of Evolutionary Biologists, Satellite Meeting (virtual)  
 Biology Seminar, University of Nebraska, Lincoln (virtual)  
 Statistics Seminar, Washington University in St. Louis (virtual)
- 2020 Biology Seminar, University of Missouri, St. Louis (virtual)

2019	Living Earth Collaborative Seminar, Washington University in St. Louis (virtual) St. Louis Ecology, Evolution & Conservation Seminar, Lewis & Clark Community College Biological Sciences Seminar, Auburn University
2018	Plant Biology Seminar, University of Georgia Evolution & Systematics Seminar, University of Connecticut Phyloseminar, an online society-sponsored seminar ( <a href="http://phyloseminar.org/">http://phyloseminar.org/</a> ) Department of Biology Seminar, University of Oregon Department of Biology Seminar, Washington University in St. Louis
2017	Symposium on Computational Paleobiology, Geological Society of America, Seattle Systematics Seminar, Swedish Museum of Natural History
2016	Computational Genomics Seminar, Temple University
2015	Symposium on Parametric Biogeography, Evolution Conference in Guarujá, Brazil
2013	Phylogenetics & Evolutionary Biology Seminar, North Carolina State University Workshop on Mathematics for an Evolving Biodiversity, University of Montréal Center for Population Genomics Seminar, UC Davis

#### **Presentations – Contributed**

2022	International Biogeography Society Conference (virtual)
2022	Evolution Conference (virtual)
2021	Evolution Conference (virtual)
2020	Midcontinent Paleobotanical Colloquium (virtual)
2019	Evolution Conference in Providence Yale Institute for Biospheric Studies Seminar
2017	External Advisory Board Meeting, Yale Institute for Biospheric Studies Evolution Conference in Portland
2016	Ernst Mayr Symposium, Evolution Conference, Austin
2013	Evolution Conference in Snowbird Center for Theoretical Evolutionary Genomics, UC Berkeley
2012	Center for Theoretical Evolutionary Genomics, UC Berkeley Evolution Conference in Ottawa

#### **Professional Reviews**

Journals	American Journal of Botany, Annals of Botany, Bioinformatics, BMC Evolutionary Biology, Evolution, Genome Biology & Evolution, Journal of Biogeography, Methods in Evolution & Ecology, Methods in Ecology & Evolution, Molecular Biology & Evolution, Molecular Phylogenetics & Evolution, Nature Communications, New Phytologist, Paleobiology, Proceedings of the Royal Society B, Proceedings of the National Academy of Sciences USA, Systematic Biology, Trends in Ecology & Evolution, Zoological Journal of the Linnean Society
Grants	NSF reviewer, SSB Ernst Mayr applications, SSB graduate award grants

#### **University Service**

2022	Hiring Committee for Genetics & Genomics Faculty
2022	Hiring Committee for Urban Biology & Environmental Justice Faculty
2021—	Faculty mentor for WUSTL Chapter within the Society for Advancement of Chicanos/Hispanics and Native Americans in Science
2019—	WUSTL Biology Curriculum Committee
2019—	EEPB Steering Committee
2019—	EEPB Admissions Committee
2012—2013	Co-chair, Integrative Biology Graduate Student Assembly, UC Berkeley

**Society Service**

2022—	Associate Editor for Systematic Biology
2021—	Legacy Committee member for Society for Systematic Biologists
2021,2022	One-off ERC mentor for Evolution meeting
2019—2022	Council member for Society for Systematic Biologists
2018—2022	Editorial board member for Systematic Biology
2019	Symposium organizer for Society of Systematic Biologists on The Bright Side of Phylogenetics

**Teaching & Courses**

2020—2022	Instructor, BIOL4220, Practical Bioinformatics, WUSTL
2022	Invited Lecturer, EEB103, Statistical Phylogenetics & Speciation, UC Davis
2021	Instructor, BIOL580, EEPB Graduate Seminar, WUSTL
2016	Invited Lecturer, BIOL1425, Phylogenetic Biology, Brown University
2014	Invited Lecturer, IB87, Bioinformatics, UC Berkeley
2013	Invited Lecturer, EEB101, Macroevolution, UC Davis

**Teaching Assistance**

2012	Teaching Assistant, IB200A, Principles of Phylogenetics, UC Berkeley
2011	Teaching Assistant, IB164, Human Genetics and Genomics, UC Berkeley

**Research Lab Mentoring & Advising***WUSTL Postdoctoral Scholar*

2021—	Fábio Mendes, Statistical Phylogenetics
2019—2021	Mariana Braga, Evolution & Ecology

*ORISE Postdoctoral Scholar*

2021—	Ammon Thompson, Statistical Epidemiology
-------	--

*WUSTL PhD Student (advised)*

2021—	Sarah Swiston, EEPB
2022—	Sean McHugh, EEPB

*WUSTL Undergraduate*

2022—	Yu (Sunny) Zichen, Mathematics
2021—	Mihir Shah, Biomedical Engineering
2021—	Walker Sexton, Biology
2021	Ernie Ramos, Mathematics

*UC Berkeley Undergraduate*

2013—2015	Bryan Wang, Mathematics
2012—2014	Jaya Narasimhan, Computer Science

**PhD Program Mentoring & Advising***WUSTL PhD Student (rotation)*

2022	Preston Pennington, EEPB
2020	Justin Baldwin, EEPB
2020	Aryeh Miller, EEPB

*WUSTL dissertation committee member*

2022—	Justin Baldwin, EEPB
2022—	Aryeh Miller, EEPB
2022—	Changxu Fan, Immunology
2022—	Jenna Lin, BBSB
2021—	Brock Mashburn, EEPB
2021—	Jhan Salazar, EEPB
2020—	Wen-His Kuo, EEPB
2020—2022	Erika Schumacher, EEPB
2020	Qi Wang, Statistics

### **Outreach**

2021	Instructor, Stay-at-Home RevBayes Workshop (virtual)
2020	Instructor, Stay-at-Home RevBayes Workshop (virtual)
2018—2019	Instructor, Molecular Evolution Workshop, Woods Hole
2019	Instructor, Bodega Phylogenetics Workshop, UC Davis
2019	Instructor, RevBayes Workshop, Yale University, New Haven
2017	Instructor, Fossil tip-dating with RevBayes, GSA Meeting, Seattle
2017	Instructor, Biogeography with RevBayes, SSB Meeting, Baton Rouge
2017	Instructor, Introduction to RevBayes, Yale, New Haven
2014—2016	Teaching Assistant, Molecular Evolution Workshop, Woods Hole
2015	Instructor, RevBayes Workshop, UC Berkeley
2014, 2015	Instructor, Bodega Phylogenetics Workshop, UC Davis
2014	Instructor, Applied Phylogenetics Workshop, NESCent
2014, 2015	Guest Lecturer, Berkeley High School Science Outreach
2014	Guest Lecturer, Bay Area Scientists in Schools
2009—2010	Tutor, 826 Valencia volunteer for English Language Learners

### **Professional Societies**

2009—	Society of Systematic Biologists
2012—	Society for the Study of Evolution
2018—	International Biogeography Society
2021—	Society for Advancement of Chicanos/Hispanics and Native Americans in Science
2017—2020	Geological Society of America

### **Professional Experience**

2005—2008	IT Consultant, Varsity Technologies, San Francisco, CA
-----------	--

### **Advisors**

PhD	John P. Huelsenbeck Department of Integrative Biology UC Berkeley
Postdoctoral	Tracy A. Heath Department of Ecology, Evolution, and Organismal Biology Iowa State University
Postdoctoral	Michael J. Donoghue Department of Ecology and Evolutionary Biology Yale University