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Education

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- 2014** Ph.D., Plant Molecular and Cellular Biology
University of Florida, Gainesville, Florida
- 2010** B.S. (*Summa cum laude*), Botany
Department of Biology
University of Florida, Gainesville, Florida

Work Experience

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- 2024-Current Lead Research Analyst, Ag Data Services, Purdue University (2024 – Current)

Research Experience

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- 2018-2024 Post-Doctoral Fellow, Biochemistry Department, Purdue University
- 2014-2018 Post-Doctoral Fellow, Department of Human Genetics, University of Michigan
- 2010-2014 Ph.D. Student, Plant Molecular and Cellular Biology, University of Florida

Publications

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- Timothy R. Fallon, Vikram V. Shende, Igor H. Wierzbicki, **Amanda L. Pendleton**, Nathan F. Watervoot, Robert P. Auber, David J. Gonzalez, Jennifer H. Wisecaver, Bradley S. Moore. (2024) Giant polyketide synthase enzymes in the biosynthesis of giant marine polyether toxins. *Science*. (In Review).
<https://doi.org/10.1101/2024.01.29.577497>
- Katharine E. Eastman, **Amanda L. Pendleton**, Mearaj A. Shaikh, Thiti Suttiyut, Raeya Ogas, Paxton Tomko, Gregory Gavelis, Joshua R. Widhalm, Jennifer H. Wisecaver. (2023) “A reference genome for the long-term kleptoplast-retaining sea slug *Elysia crispata* morphotype clarki” *G3: Genes, Genomes, Genetics* 13(12): jkad234. <https://doi.org/10.1093/g3journal/jkad234>
- Abigail S. Jarosz, **Amanda L. Pendleton**, Michael J. Lashbrook, Erica Cech, Madison Altieri, Austin Kunch, Jaime F. Modiano, Julia V. Halo. (2023) “Expression and high levels of insertional polymorphism of an endogenous gammaretrovirus lineage in dogs” *PLoS Genetics* 19(12): e1011083.
<https://doi.org/10.1371/journal.pgen.1011083>
- Jennifer H. Wisecaver, Robert P. Auber, **Amanda L. Pendleton**, Nathan F. Watervoot*, Timothy R. Fallon, Olivia Riedling, Bradley S. Moore, William W. Driscoll. Extreme genome diversity and cryptic speciation in a harmful algal bloom forming eukaryote. *Current Biology*. 33(11) <https://doi.org/10.1016/j.cub.2023.05.003>
- Julia V. Halo*, **Amanda L. Pendleton***, Feichen Shen*, Aurelien Doucet, Thomas Derrien, Christophe Hitte, Laura E. Kirby, Bridget Myers, Elzbieta Sliwerska, Sarah Emery, John V. Moran, Adam R. Boyko, Jeffrey M. Kidd. “Long-read assembly of a Great Dane genome highlights the contributions of GC-rich sequence and mobile elements to canine genomic diversity” *Proceedings of the National Academy of Sciences* (2021) 118(11). <https://doi.org/10.1073/pnas.2016274118>
* Authors contributed equally to this work.
- Peter Thielen*, **Amanda L. Pendleton***, Robert Player, Kenneth Bowden, Thomas Lawton, Jennifer H. Wisecaver. “Reference genome for the highly transformable *Setaria viridis* ME034V” *G3: Genes, Genomes, Genetics* (2020) 10(10): 3467-3478. <https://doi.org/10.1534/g3.120.401345>
* Authors contributed equally to this work.

Amanda L. Pendleton, Ph.D.

- Jing Li, Zhenxin Fan, Feichen Shen, **Amanda L. Pendleton**, Yang Song, Bisong Yue, Jeffrey M. Kidd, and Jing Li. "Genome-wide CNV study of nine *Macaca* species reveals their genetic divergence and evolutionary adaptations" (2020) *Genome Biology and Evolution*. <https://doi.org/10.1093/gbe/evaa200>
- Andrew G. DeMarco, Kedric Milholland, **Amanda L. Pendleton**, Daniel Wesenberg, Monessha Nambiar, Antonella Pepe, Stefan Paula, Jean Chmielewski, Jennifer H. Wisecaver, W. Andy Tao, Mark C. Hall "Conservation of Cdc14 phosphatase specificity in plant fungal pathogens: implications for antifungal development" (2020) *Scientific Reports* 10:2073. <https://doi.org/10.1038/s41598-020-68921-3>
- Robert P. Auber, Thiti Suttiyut, Rachel M. McCoy, Manoj Ghaste, Joseph W. Crook, **Amanda L. Pendleton**, Joshua R. Widhalm, and Jennifer H. Wisecaver (2020) "Hybrid *de novo* genome assembly of red gromwell (*Lithospermum erythrorhizon* Siebold & Zucc.) reveals evolutionary insight into shikonin biosynthesis" *Horticultural Research* 7:82. <https://doi.org/10.1038/s41438-020-0301-9> (PMID: 32528694)
- Julia V. Halo, **Amanda L. Pendleton**, Abigail S. Jarosz, Robert J. Gifford, Malika L. Day, Jeffrey Kidd (2019) "Origin and recent expansion of an endogenous gammaretroviral lineage in canids" *Retrovirology*. 16(1): 1-25. <https://doi.org/10.1186/s12977-019-0468-z>
- Amanda L. Pendleton**, Feichen Shen, Angela M. Taravella, Sarah Emery, Krishna R. Veeramah, Adam R. Boyko and Jeffrey M. Kidd. (2018) "Comparison of village dog and wolf genomes highlights the role of the neural crest in dog domestication" *BMC Biology* 16(1):1-21. <https://doi.org/10.1186/s12915-018-0535-2>
- Laura Botigue*, Shiya Song*, Amelie Scheu*, Shyamalika Gopalan, **Amanda L. Pendleton**, Matthew Oetjens, Angela Taravella, Timo Seregély, Andrea Zeeb-Lanz, Rose-Marie Arbogast, Dean Bobo, Martina Unterländer, Kevin Daly, Jeffrey Kidd, Joachim Burger, Krishna Veeramah. (2017) "Genomic analysis of Neolithic dogs from Central Europe" *Nature Communications* 8:1-11. <https://doi.org/10.1038/ncomms16082>
- *Authors contributed equally to this work.
- Amanda L. Pendleton**, Katherine E. Smith, Nicolas Feu, Francis Martin, Igor Grigoriev, Richard Hamelin, C. Dana Nelson, J. Gordon Burleigh, John M. Davis. (2014) "Patterns of duplications and losses in gene families of basidiomycetes and rust pathogens" *Frontiers in Plant Science*. <https://doi.org/10.3389/fpls.2014.00299>

Selected Funding and Awards

2022	Purdue AgSEED Grant (Co-Investigator): \$48,853 <u>Co-Investigators:</u> Maria Soledad Sepulveda (PI), Amanda Pendleton , Tyler Hoskins, Jason Cannon
2019	Purdue Summer Course Design Fellowship: \$1,000
2015-2017	NHGRI Genome Science Training Program Fellowship: \$87,000