

# Aryeh H. Miller

Computational Biologist

✉ aryehmiller@wustl.edu

☎ 314-935-6860

📍 McDonnell Hall 415, Washington University

🆔 0000-0002-5031-3991

🔄 AryehMiller

🌐 <https://aryehmiller.github.io/>

## Education

|                     |   |
|---------------------|---|
| Aug 2020 – Present  | <b>School of Medicine, Washington University in St. Louis</b><br><i>PhD in Ecology and Evolutionary Biology</i> |
| Aug 2016 – May 2020 | <b>University of North Carolina at Asheville</b><br><i>BS in Biology with honors</i>                            |

## Experience

|             |   |
|-------------|---|
| 2018 - 2020 | <b>Research Fellow – National Museum of Natural History, Smithsonian Institution</b> <ul style="list-style-type: none"><li>Designed, deployed, and executed a cost-efficient, high-throughput wet and dry lab pipeline to generate multi-locus genetic sequence data of vertebrate biodiversity in Vietnam.</li></ul> |
|-------------|---|

## Technical Expertise

### Computing

- Programming, data manipulation and visualization in R and Python, including use of popular packages and libraries (e.g., Tidyverse, pandas).
- Statistical expertise in maximum likelihood estimation and Bayesian inference approaches, Markov models, multivariate and machine learning techniques (random forest, K-nearest neighbor, K-means clustering, decision trees, dimensionality reduction [PCA, LDA]).
- Genomic data analysis expertise in methods for whole-genome sequencing and assembly, variant calling (e.g., Trimmomatic, BWA, samtools, GATK, vcftools, bcftools), and phylogenomic inference (e.g., IQ-TREE, RevBayes, RAxML, SVDQuartets), as well as bioinformatic workflow management and engineering (Snakemake).
- High-performance computing (HPC) environment proficiency using bash, SLURM scripting, and docker images.

### Wet-lab

- Experienced with laboratory genetics, including DNA extraction manual kits and the AutoGen® system, PCR, clean-up, gel electrophoresis, Sanger sequencing, and DNA quantitation with Qubit® 3.0.

### Relevant Formal Coursework

- Big Data Science, Genomics, Bio-statistics, Population Genetics, Evolution, Cell Biology

## Leadership and Teaching Experience

|                 |  |
|-----------------|--|
| Fall 2021, 2023 | <b>Teaching Assistant (Bio 3501: Evolution), Washington University in St. Louis</b> <ul style="list-style-type: none"><li>Led weekly discussion sections where students would grapple with recent peer-reviewed articles in the evolutionary biology literature, graded exams and essays, and held weekly office hours and several exam review sessions to teach students.</li></ul> |
| 2021-Present    | <b>Mentor and Program Leader, The Young Scientist Program</b> <ul style="list-style-type: none"><li>Cultivating mentorship and teaching competencies by engaging St. Louis public high school students to hone their own knowledge of biology by conducting supervised original research and explore careers in STEM.</li></ul>  |

## Selected Recent Honors and Awards

1. National Science Foundation Graduate Research Fellow Award Finalist (2020, 2021)

- 2. University Research Scholar, UNC Asheville (2020)
- 3. Bernhardt-Perry Award for Undergraduate Research, UNC Asheville (2020)
- 4. Smithsonian Institution Global Genome Initiative Award (2018)

Highlighted Publications

---

|      |    |  |
|------|----|--|
| 2023 | 1. | <b>Miller, A. H.</b> , Stroud, J. T. & Losos, J. B. The ecology and evolution of key innovations. <i>Trends in Ecology and Evolution</i> <b>38</b> , 122–131 (2023).   |
| 2022 | 2. | <b>Miller, A. H.</b> & Stroud, J. T. Novel tests of the key innovation hypothesis: Adhesive toepads in arboreal lizards. <i>Systematic Biology</i> <b>71</b> , 139–152 (2022).   |
|      | 3. | Reynolds, R. G. <i>et al.</i> Phylogenomics and Historical Biogeography of West Indian Rock Iguanas (genus <i>Cyclura</i> ). <i>Molecular Phylogenetics and Evolution</i> <b>174</b> , 107548 (2022).  |
| 2020 | 4. | <b>Miller, A. H.</b> <i>et al.</i> Discovery of a new species of enigmatic odd-scaled snake (Serpentes: Xenodermidae: Achalinus) from Ha Giang Province, Vietnam. <i>Copeia</i> <b>108</b> , 796–808 (2020).                                 |
| 2019 | 5. | <b>Miller, A. H.</b> <i>et al.</i> The complete mitochondrial genome of the critically endangered Lesser Antillean iguana ( <i>Iguana delicatissima</i> ; Squamata: Iguanidae). <i>Mitochondrial DNA Part B</i> <b>4</b> , 2479–2481 (2019). |