Email: <a href="mailto:aw3746@cumc.columbia.edu">aw3746@cumc.columbia.edu</a>
Website: <a href="mailto:www.amandaewilson.com">www.amandaewilson.com</a>

GitHub: github.com/aewilson96

### **Research Experience**

Postdoctoral Research Scientist June 2024 - Present

Dr. Laura Landweber's Research Group

Department of Biochemistry and Molecular Biophysics

Columbia University, New York, NY

Modeling rearrangement of ciliate genomes

Research Assistant 2018 - 2024

Dr. David A. Liberles' Research Group

Department of Biology

Temple University, Philadelphia, PA

Modeling evolutionary processes of duplicated genes

Research Assistant 2017

Dr. Robert Minckley's Research Group

Department of Biology

University of Rochester, Rochester, NY

Evolution and ecology of bees and their floral hosts

### **Education**

Ph.D. in Biology Awarded May 2024

Temple University, Philadelphia, PA

Dissertation: <u>SELECTIVE FORCES SHAPING DUPLICATE GENE EVOLUTION</u>: <u>INSIGHTS FROM STOCHASTIC MODELING AND PATTERNS OF RETENTION</u>

*Advisor:* Professor David Liberles

Committee: Professor Sergei Pond, Professor Rachel Spigler,

Professor Anne-Ruxandra Carvunis

GPA: 3.95

B.S. in Biological Sciences

University of Rochester, Rochester, NY

Evolutionary Biology and Ecology Track

Clusters (minors): "Public Health" and "The Performing Musician"

#### **Publications**

Soewongsono AC, Diao J, Stark T, <u>Wilson AE</u>, Holland B, Liberles DA, O'Reilly MM. (2023) Matrix-analytic methods for the evolution of species trees, gene trees, and their reconciliation. arXiv [q-bio.PE]. (Under Review) https://doi.org/10.48550/arXiv.2309.06447

Assis R, Conant G, Holland B, Liberles DA, O'Reilly MM, <u>Wilson AE</u>. (2023) Models for the Retention of Duplicate Genes and Their Biological Underpinnings. *F1000Research*, 12: 1400. <a href="https://doi.org/10.12688/f1000research.141786.1">https://doi.org/10.12688/f1000research.141786.1</a>

<u>Wilson AE</u>, Liberles DA. (2023) Expectations of Duplicate Gene Retention Under the Gene Duplicability Hypothesis. *BMC Ecology and Evolution*. *BMC Ecology and Evolution*, 23: 76. https://doi.org/10.1186/s12862-023-02174-2

Wilson AE, Liberles DA. (2023) Dosage Balance as a time-dependent selective barrier to

Awarded May 2018

subfunctionalization of expression states. *BMC Ecology and Evolution,* 23: 14. https://doi.org/10.1186/s12862-023-02116-y

- Henry CN, Piper K, <u>Wilson AE</u>, Miraszek JL, Probst CS, Rong Y, Liberles DA. (2022) WGDTree: A Phylogenetic Software Tool to Examine Conditional Probabilities of Retention Following Whole Genome Duplication Events. *BMC Bioinformatics*, 23: 505. <a href="https://doi.org/10.1186/s12859-022-05042-w">https://doi.org/10.1186/s12859-022-05042-w</a>
- Anthony-Regnitz CM, <u>Wilson AE</u>, Sweazea KL, Braun EJ. (2020) Fewer Exposed Lysine Residues May Explain Relative Resistance of Chicken Serum Albumin to In Vitro Protein Glycation in Comparison to Bovine Serum Albumin. *Journal of Molecular Evolution*, 88: 653-661. https://doi.org/10.1007/s00239-020-09964-y
- <u>Wilson AE</u>, Kosater WM, Liberles DA. (2020) Evolutionary Processes and Biophysical Mechanisms: Revisiting Why Evolved Proteins Are Marginally Stable. *Journal of Molecular Evolution*, 88:415-417. <u>https://doi.org/10.1007/s00239-020-09948-v</u>

### **Conference Talks**

<u>Wilson AE</u> , Liberles DA. "Expectations of Duplicate Gene Retention Under the Gene Duplicability Hypothesis". The JME Advances Meeting 2023, Washington, D.C., USA	March 2023
<u>Wilson AE</u> , Liberles DA. "Dosage Balance as a time-dependent selective barrier to subfunctionalization of expression states". GLAM-Evogen 2022, Buffalo, NY, USA	Aug 2022
<u>Wilson AE</u> . "Modeling Whole Genome Duplication". GLAM-Evogen 2021,Virtual Conference	Aug 2021
<u>Wilson AE</u> . "Modeling probabilities of retention of gene copies after consecutive whole genome duplication events".  T3 (TelAviv, Toronto, Temple) 2021, Virtual Conference	Apr 2021
<u>Wilson AE</u> . "Modeling Probabilities of Retention of Duplicated Genes". iGem 2021, Philadelphia, PA, USA - Virtual Conference	Apr 2021
<u>Wilson AE</u> . "Modeling Probabilities of Retention of Gene Copies after Consecutive Whole Genome Duplication Events". GLAM-Evogen 2020, Rochester, NY, USA - Virtual Conference	Jul 2020
<u>Wilson AE</u> . "Modeling Probabilities of Retention of Gene Copies after Consecutive Whole Genome Duplication Events". Phylomania 2019, Hobart, TAS, Australia	Nov 2019
Conference Posters	
<u>Wilson AE</u> , Liberles DA. "Dosage Balance as a time-dependent selective barrier to subfunctionalization of expression states".  GLAM-Evogen 2022, Buffalo, NY, USA	Aug 2022
<u>Wilson AE</u> , Liberles DA. "Modeling Probabilities of Retention of Gene Copies after Consecutive Whole Genome Duplication Events".  SMBE 2021, Virtual Conference	Jul 2021
<u>Wilson AE</u> , Miraszek J, Liberles DA. <i>"Whole Genome Duplication Events drive diversification of the myostatin pathway in Salmonidae".</i> Evolution in Philadelphia Conference (EPiC) 2019, Philadelphia, PA, USA	Sep 2019

# **Teaching Experience**

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Department of Biology at Temple University:	
Intro to Organismal Biology (BIOL1111), Laboratory Teaching Assistant	Spring 2020
Intro to Cell and Molecular Biology (BIOL2112), Laboratory Teaching Assistant	Fall 2019
General Biology II (BIOL1012), Laboratory Teaching Assistant	Spring 2019
General Biology I (BIOL1011), Laboratory Teaching Assistant	Fall 2018
Department of Biology at University of Rochester:	
Animal Behavior (BIO260), Teaching Assistant	Fall 2017
Introductory Biology Lab II (BIO117P), Teaching Assistant	Summer 2017
Department of Chemistry at University of Rochester:	
Organic Chemistry II: Lab (CHM208), Teaching Assistant	Spring 2017
Organic Chemistry I: Lab (CHM207), Teaching Assistant	Fall 2016
Scientific Involvement	
Journal of Molecular Evolution: Social Media Coordinator Compose and post summaries of new publications on social media Write blog posts on Nature Ecology and Evolution Community page	2022 - 2023
Center for Computational Genetics and Genomics (CCGG): Journal Club  Attend weekly Journal Club meetings; Present a recent publication each semester	2018 - 2024
Temple University: Bioinformatics Studio	2019-2024
Attend regular meetings to engage in collaborative learning of bioinformatics pipelines	

# **Scientific Journal Peer Review**

G3; Journal of Molecular Evolution; PLOS Computational Biology; Molecular Biology and Evolution; BMC Ecology and Evolution

# **Professional Society Membership**

Society for Molecular Biology and Evolution (SMBE)

# **Academic Honors**

Academic Honors			
Awarded Doctoral Dissertation Completion Grant, Temple University	Spring 2024		
Scholarship, Prince Street Merit, University of Rochester	2014 - 2018		
National Honors Society	2010 - 2014		
Leadership and Community Service			
University of Rochester Community Symphony Orchestra	2015 - 2018		
Cellist			
Eye to Eye - Rochester Chapter	2015 - 2018		
Mentor middle school students with learning disabilities			
Leadership Roles: Secretary and Publicity Chair			