Maike L Morrison maikem@stanford.edu | MaikeMorrison.com Education 2020 -Stanford University

Stanford, CA

2016-2020 The University of Texas at Austin

PhD in Biology (Ecology & Evolution)

Austin, TX

BS in Mathematics, Dean's Scholars Honors Program

Research Experience

Sep $2020-$	Graduate Research Assistant	Stanford, CA
Present	Supervised by Dr. Noah Rosenberg in the Stanford University Department of	Biology.
Jun 2022– July 2022	Complex Systems Summer School Student Participant in the Santa Fe Institute Complex Systems Summer School.	Santa Fe, NM
Aug 2017– May 2021	Mathematical Epidemiology Undergraduate Research Assistant Supervised by Dr. Lauren Ancel Meyers in the UT Austin Department of Inte	Austin, TX

Supervised by Dr. Lauren Ancel Meyers in the UT Austin Department of Integrative Biology. May 2021 Dec 2017-**Evolutionary Genetics Undergraduate Research Assistant**

Aug 2020 Supervised by Dr. Mark Kirkpatrick in the UT Austin Department of Integrative Biology.

Senior Honors Mathematics Thesis Aug 2019-

Austin, TX

May 2020 Supervised by Dr. Stephen Walker in the UT Austin Departments of Mathematics.

May 2019-UC San Francisco Amgen Scholars Program

San Francsico, CA

Jul 2019 Supervised by Dr. John Witte in the UCSF Department of Biostatistics & Epidemiology. Participant in UCSF's Summer Research Training Program.

Jun 2018-The Mathematical Biosciences Institute NSF REU State College, PA

Supervised by Dr. Ephraim Hanks in the Penn State University Department of Statistics. Par-Aug 2018 ticipant in the MBI Research Experience for Undergraduates program at Ohio State University.

Peer-Reviewed Publications

Morrison ML, Alcala N, Rosenberg NA. FSTRUCT: an F_{ST}-based tool for measuring ancestry variation in inference of population structure. Molecular Ecology Resources, 22, 2614–2626.

Ingle T, Morrison ML, Wang X, Mercer T, Karman V, Fox S, Meyers LA (2021). Projecting COVID-19 isolation bed requirements for people experiencing homelessness. PLOS One, 16(5), e0251153.

Kachuri L*, Francis SS*, Morrison ML, Wendt GA, Bossé Y, Cavazos TB, Rashkin SR, Ziv E, Witte JS (2020). The landscape of host genetic factors involved in immune response to common viral infections. Genome Medicine 12(93): 1-18.

Morrison M*, Castro LA*, Meyers LA (2020). Conscientious vaccination exemptions in kindergarten to eighth-grade children across Texas schools from 2012 to 2018: A regression analysis. PLOS Medicine 17(3): e1003049.

^{*}These authors contributed equally

Honors, Awards, & Fellowships			
2022	Stanford Biology Department Excellence in Teaching Award		
2021 – 2025	NSF Graduate Research Fellowship Program		
2020 – 2025	Stanford Graduate Fellowship		
2020	University Co-op Mitchell Award for Outstanding Undergraduate Research		
2020	Dean's Honored Graduate: University of Texas College of Natural Sciences		
2020	UT Austin Mathematics Special Departmental Honors		
2020	NSF Graduate Research Fellowship Program, Honorable Mention		
2019–2020	Alan Kaylor Cline Dean's Scholars Scholarship		
2019–2020	Phi Beta Kappa, junior year election		
2019	Undergraduate Diversity at Evolution Award		
2019	Barry M. Goldwater Scholarship		
2018 – 2020	University of Texas University Honors: College Scholar		
2017	Freshman Research Initiative Summer Research Fellowship		
2016-2020	University of Texas College of Natural Sciences Scholarship		
2016-2020	Dean's Scholars Honors Program		
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Winter 2022	BIO 187: Mathematical Population Biology Graduate Teaching Assistant	Stanford University Dr. Noah Rosenberg	
Winter	BIO 187: Mathematical Population Biology Graduate Teaching Assistant BIO 85: Evolution	·	
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Conference Presentations				
Apr 2019	Undergraduate Research Forum at UT Austin "How Many Species Make Up the World's Most Abundant Photosynthetic Organism?" FSTI Award for Excellence in Oral Presentation			
Nov 2018	Capital of Texas Undergraduate Research Conference "Analyzing Conscientious Vaccination Exemption Rates in Texas." Award of Excellence for an Outstanding Oral Presentation of Research			
Aug 2018	Mathematical Biosciences Institute Capstone Conference Columbus, OH "Distinguishing Resource Selection from Heavy-Tailed Dispersal in Spatial Epidemic Models." <u>Link to recording.</u> (co-presented with Emily Strong)			
Poster Pr	<u>esentations</u>			
June 2022	Population, Evolutionary, & Quantitative Genetics Conference $^{\circ}$ Pacific Grove, CA "FSTruct: An F_{ST} -based tool for quantifying ancestry variability." $^{\circ}$ Genetics Society of America Poster Award			
Jun 2019	Evolution Meeting "How Many Ecotypes are Comprised by the World's Most Abundant Photosynthetic Organism?" Undergraduate Diversity at Evolution Award Program			
Apr 2019	Longhorn Research Poster Session "Modeling Conscientious Vaccination Rates in Texas Metropolitan Areas." Austin, TX			
Aug 2018	Mathematical Biosciences Institute Capstone Conference Columbus, OH "Distinguishing Resource Selection from Heavy-Tailed Dispersal in Spatial Epidemic Models."			
Apr 2018	Undergraduate Research Forum at the University of Texas at Austin Austin, TX "Conscientious Vaccination Exemption Rates in Texas: A Longitudinal Analysis of Trends and Predictors."			