# JENNIFER BLANC

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### **EDUCATION**

# University of Chicago

September 2018 - September 2024 (expected)

Ph.D in Human Genetics

• NHGRI F31 Ruth L. Kirschstein Predoctoral Individual National Research Service Award

# University of California, Davis

October 2014 - June 2018

B.S in Genetics and Genomics, minor in Statistics

- Graduated with highest honors (top 2%)
- Top Senior in Genetics and Genomics

#### RESEARCH EXPERIENCE

PhD Candidate September 2018 - Present

Department of Human Genetics, University of Chicago

- Studies human adaptation and the evolution of complex traits in Dr. Jeremy Berg's lab
- Developed novel statistical approach to control for confounding in polygenic score analyses

#### Student Researcher

September 2017 - August 2018

Department of Evolution and Ecology, University of California, Davis

- Completed undergraduate honors thesis in the population genetics lab of Dr. Graham Coop
- Worked with Dr. Emily Josephs on detecting local adaptation in Maize using gene expression data

# Laboratory Intern

June 2017 - August 2017

Icahn School of Medicine at Mount Sinai

- Summer Undergraduate Research Program intern for Dr. Eimear Kenny
- Studied the demographic history of Oceanic populations using identity by descent

Student Researcher April 2015 - June 2017

Department of Evolution and Ecology, University of California, Davis

• Lead independent project mapping the genetic basis of cytoplasmic incompatibility in *D. melanogaster* in Dr. Michael Turelli's lab

#### **High School Intern**

April 2015 - June 2015

Department of Plant Sciences, University of California, Davis

- Learned basic laboratory techniques including PCR and bacterial cloning
- Assisted on a project to sequence the genes involved in the first step of a hydrolysable tannin pathway in the lab of Dr. Li Tian

# TECHNIAL SKILLS

**Programming** R (including tidyverse and base R), Python, shell scripting

Data Analysis Snakemake, Plink, GCTA, SAMtools, GATK

**Laboratory** PCR, DNA Extraction/Purification, qPCR, Inverse PCR, Bacterial Cloning,

Gel Electrophoresis, *Drosophila* stock keeping and crossing experiments

#### **PUBLICATIONS**

Blanc, Jennifer, and Jeremy J. Berg. "Testing for differences in polygenic scores in the presence of confounding." bioRxiv (2023): 2023-03. https://doi.org/10.1101/2023.03.12.532301

**Blanc, Jennifer** and Jeremy J. Berg. 2020. "Polygenic Scores: How Well Can We Separate Genetics from the Environment?" eLife 9: e64948. https://doi.org/10.7554/eLife.64948

Blanc, Jennifer, Karl AG Kremling, Edward Buckler, and Emily B. Josephs. "Local adaptation contributes to gene expression divergence in maize." G3 11, no. 2 (2021): jkab004. https://doi.org/10.1093/g3journal/jkab004.

#### TEACHING EXPERIENCE

### Course Instructor

QBio: Quantitative Analysis Bootcamp - Population Genetics Workshop September 2022 Taught two sections of a population genetics workshop to first-year graduate students as part of a week long bootcamp focused on establishing familiarity with computational biology and analysis in R.

# Citizen Science at Bard College

January 2022 - February 2022

Taught 3 week intensive, project-based undergraduate course on scientific literacy and data science.

# Teaching Assistant

#### Teaching Assistant: Human Variation and Disease

January 2022 - March 2022

University of Chicago Graduate Course

#### Teaching Assistant: QBio (Quantitative Analysis Bootcamp)

September 2021

University of Chicago Graduate Student Bootcamp

# Teaching Assistant: Human Genetics 1

September 2020 - December 2020

University of Chicago Graduate Course

#### Teaching Assistant: Human Variation and Disease

March 2020 - June 2020

University of Chicago Graduate Course

Awarded Divisional Teaching Assistantship Award in the Biological Sciences

Summer Abroad Tutor: Intro to Genes and Gene Expression August 2016 - September 2016 UC Davis Undergraduate Course

#### **PRESENTATIONS**

#### Invited Speaker

### Adaptation in Structured Populations

June 2023

Department of Human Genetics the Quantitative Life Sciences Program at McGill University (virtual) Testing for differences polygenic scores in the presence of confounding

#### DISCOVER: What you can do with a career in natural or social sciences

April 2023

Saint Norbert's College

**Evolution of Complex Traits** 

# Find Yourself: Tracing Human Origins Using DNA

April 2021

New York Academy of Sciences course for high school students (virtual) Guaranteeing unbiasedness in selection tests based on polygenic scores

# Conference Oral Presentations

American Society of Human Genetics Annual Meeting Testing for differences in polygenic scores in the presence of confounding	November 2023
Probabilistic Modeling in Genomics Analyzing the role of population structure in polygenic score analyses	March 2023
Population, Evolutionary, and Quantitative Genetics Conference Guaranteeing unbiasedness in selection tests based on polygenic scores	June 2022
Probabilistic Modeling in Genomics Guaranteeing unbiasedness in selection tests based on polygenic scores	March 2022
Midwest Population Genetics VII Guaranteeing unbiasedness in selection tests based on polygenic scores	August 2021
UC Davis Undergraduate Research Conference Detecting Local adaptation in gene expression in maize	April 2018
Conference Poster Presentations	
The Allied Genetics Conference: PEQG Testing for differences in polygenic scores in the presence of confounding	March 2024
Midwest Population Genetics VIII Testing for differences in polygenic scores in the presence of confounding	August 2023
The Mitchell Conference on Human Genetics Testing for differences in polygenic scores in the presence of confounding	May 2023
American Society of Human Genetics Annual Meeting Guaranteeing unbiasedness in selection tests based on polygenic scores	October 2022
American Society of Human Genetics Annual Meeting Guaranteeing unbiasedness in selection tests based on polygenic scores	October 2021
Probabilistic Modeling in Genomics Guaranteeing unbiasedness in tests of polygenic adaptation Reviewer's Choice selected abstract (top 10%)	April 2021
American Society of Human Genetics Annual Meeting Using derived allele status to detect and correct for stratification in GWAS summar	October 2020 y statistics
Midwest Population Genetics VI Detecting local adaptation in gene expression in maize Winner of best graduate student poster	August 2019
Summer Undergraduate Research Program Symposium (Mount Sinai) Reconstructing the population history of Oceania using identity by descent	August 2017
UC Davis Undergraduate Research Conference Mapping genes controlling the level of cytoplasmic incompatibility in <i>Drosophila me</i>	April 2016
CADEMIC ACHIEVEMENTS	

Awarded F31 Ruth L. Kirschstein Predoctoral Individual National Research Service Award

Semi-finalist for Provosts Office Dissertation Completion Fellowship (2023-2024)

Won Divisional Teaching Assistantship Award in the Biological Sciences Division 2019-2020

Graduated UC Davis with Highest Honors (top 2% of graduating class)

Won Outstanding Senior in Genetics and Genomics (2018). Nominated by Dr. Michael Turelli and given to the top student in each major for academic and research achievement.

Graduated from the UC Davis Honors Program and completed Honors thesis.

UC Davis Reagents Scholar (top 4% of applicants)

#### PROFESSIONAL AFFILIATIONS

Genetics Society of America (GSA)

2022-Present

American Society of Human Genetics (ASHG)

2021-Present

#### OUTREACH AND SERVICE

Department of Human Genetics Diversity, Equity, and Inclusion Committee Member

2023-Present

UC COMBO Communication and Education Chair

2022 - Present

• In charge of organizing coding workshops and marketing for UC COMBO (UChicago Computational Biology Outreach)

item Program in Computational Biology Snakemake Tutorial Development

2024

 $\bullet$  Designed an introduction to the workflow management system Snakemake

Student Representative, Human Genetics Graduate Program

2019-2021

• Organized graduate recruitment events, the annual Molecular Biosciences retreat, and events for Human Genetics graduate students

Software Carpentry Instructor

2019 - 2021

• Led a virtual Introductory to R and Unix workshop for graduate students, postdocs, and staff in the BSD community

History of Race and Genetics Reading Group

2020

• Participated in and wrote reflections for a reading group on the history of race and genetics, helped design the website to publish materials

Computational STEM Lab Instructor

2019-2020

• Helped design and led in person and virtual workshops on computational biology in Python for Chicago area high school and middle school age students

Graduate student tutor 2019-2021

- HGEN 47400: Introduction to Probability and Statistics for Geneticists
- HGEN 47100: Statistical Genetics
- ECEV 35600: Population Genetics

Volunteer in Chicago Public Schools

2019

Volunteer at Explorit Science Center

2009-2018