

## Justin Williams, Ph.D.

### Contact

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### Software Skills

<b>R</b>	<i>Preferred</i>
<b>Git</b>	<i>GitLab, GitHub</i>
<b>Python</b>	<i>Research</i>
<b>SQL</b>	<i>PostgreSQL, MySQL</i>
<b>CI/CD</b>	<i>GitLab, GitHub</i>
<b>AWS</b>	<i>S3, EC2, ECR</i>
<b>Docker</b>	<i>Productization</i>
<b>SAS</b>	<i>Coursework</i>
<b>Stata</b>	<i>Coursework</i>

### Language

<b>English</b>	<i>Native</i>
<b>Spanish</b>	<i>Elementary</i>

### Biography

Graduated from the University of California, Los Angeles (UCLA) with a Ph.D. in Biostatistics. Currently I am employed with the Los Angeles Dodgers as a Senior Quantitative Analyst. I have a wide variety of experiences interning in industry along with extensive research work within the field of autism spectrum disorder

### Professional Experience

#### Senior Quantitative Analyst

Los Angeles Dodgers-Los Angeles, CA

11/2022 - Present

#### Quantitative Analyst

Los Angeles Dodgers-Los Angeles, CA

08/2020 - 10/2022

- Built performance model in collaboration with amateur scouting department to quantify expected hitter performance
- Led productization efforts for prediction and re-train tasks using Airflow and dependency management solutions such as `renv` (R) and `poetry` (Python)
- Developed and deployed internal repository to host proprietary R packages

#### Product Development Biostatistics Intern

Genentech-South San Francisco, CA

Summer 2019

- Designed software to simulate longitudinal differential abundance for microbiome: `microbiomeDASim`
  - Flexibly specify form of the trend over time including polynomial, oscillating, or hockey stick trends
  - Define desired sample size, number of repeated measures, and signal:noise ratio
  - Multiple choices for longitudinal dependence including:  $AR(1)$ , compound, or independent
- Compared multiple methods for estimating differential abundance over time

#### Biostatistics R&D Intern

Alcon-Ft. Worth, TX

Summer 2018

- Developed methodology for parameter estimation of censored data from truncated normal distribution
- Investigated available methods for estimation with left censoring using R and SAS
- Produced functions and macros to simulate data and calculate bias metrics
- Applied methods to estimate parameters for historical clinical trial data

## Education

09/2016 - 12/2020

### Biostatistics (Ph.D.)

University of California Los Angeles

Dissertation: "Methods for estimating causal effects for multivariate continuous exposure".

Advisor: Catherine Crespi

09/2014 - 06/2016

### Biostatistics (M.S.)

University of California Los Angeles

Bachelor's thesis: "Bayesian hierarchical spatial analysis of autism spectrum disorder services in the Los Angeles Unified School District".

Advisor: Rob Weiss

09/2009 - 05/2013

### Mathematics (B.A.)

Boston College

## Research Interests

- ▶ Causal Inference
- ▶ Longitudinal Analysis
- ▶ Bayesian Analysis
- ▶ Machine Learning

## Personal Interests

- ▶ Baseball
- ▶ Hiking
- ▶ Cooking
- ▶ Travel

### Predictive Analytics Intern

Ingram Micro-Irvine, CA

Summer 2016

- Illustrated regional product demand for products to inform warehouse stocking decisions
- Pulled purchasing and warehouse transaction information from servers via SQL
- Engineered product similarity scores based on feature list with mixed scale variables

## Research Experience

### Graduate Student Researcher

Connie Kasari Lab, UCLA Semel Institute

2015 - 2019

- Constructed Bayesian multi-level hierarchical model incorporating spatial random effects
- Developed longitudinal data visualization tools available in GitHub R package (ggplot.spaghetti)
- Clinical trial longitudinal analysis using mixed effects and generalized estimating equations
- Adjusted for empirical trends using zero-inflated and hurdle models with count outcomes
- Automated analysis for inter rater reliability
- Data management and data cleaning for multisite clinical trial database
- Co-authored multiple papers as primary statistician

## Teaching Experience

### Special Reader:

#### Computer Management of Health Data

UCLA, Biostatistics Department

Fall Quarter 2017

- Taught data management tools with SAS to 23 first-year Biostatistics graduate students
- Introduced tools for creating randomization schemes and generating reproducible data

### Special Reader:

#### Basic Biostatistics

UCLA, Biostatistics Department

Winter Quarter 2016

- Led weekly lab sections using Stata on topics such as linear regressions, ANOVA, logistic regression, and non-parametric tests
- Designed and administered discussion sections weekly
- Graded homework and lab assignments

## Awards & Honors

### Dissertation Year Fellowship (\$20,000)

#### Awarded By: UCLA Graduate Division

Received: December 2019 – December 2020

### Most Outstanding Oral Presentation (\$500)

#### Awarded By: Western North American Region of the International Biometric Society

Received: June 2019

### Juneal Marie Smith Fellowship in International Nutrition (\$2,500)

#### Awarded By: UCLA Fielding School of Public Health

Received: June 2019

### Graduate Summer Research Mentorship (\$6,000)

#### Awarded By: UCLA Graduate Division

Received: June 2017 - September 2017

## Publications

- Sturm, A., **Williams, J.**, and Kasari, C. (2021). Who gains and who loses? Sociodemographic disparities in access to special education services among autistic students. *Autism Research*, 14:1621–1632. doi:10.1002/aur.2517.
- Melamed, K.H, **Williams, J.**, Wang, X., Hu S., Nguyen C., Cui, J., & Deng, J.C. Development of secondary bacterial pneumonia in adults presenting with influenza versus noninfluenza viral respiratory infection. *Therapeutic Advances in Respiratory Diseases*, 14. doi:10.117/1753466620963026.
- **Williams, J.R.**, and Crespi, C.M. (2020). Causal inference for multiple continuous exposures via the multivariate generalized propensity score. *arXiv preprint. arXiv:2008.13767*
- **Williams, J.R.**, Kim, H. and Crespi, C.M. (2020). Modeling observations with a detection limit using a truncated normal distribution with censoring. *BMC Med Res Methodol*, 20:170. doi:10.1186/s12874-020-01032-9
- **Williams, J.**, Bravo H.C., Tom J., and Paulson J.N (2020). *microbiomeDASim*: Simulating longitudinal differential abundance for microbiome data [version 2; peer review: 2 approved]. *F1000Research*, 8:1769. doi:10.12688/f1000research.20660.2
- Dean, M., **Williams, J.**, Kasari, C. and Orlich, O. (2020). Adolescents with autism spectrum disorder and social skills groups at school: A randomized trial comparing intervention environment and peer composition. *School Psychology Review*, 49(1):60-73. doi:10.1080/2372966X.2020.1716636
- Gulsrud, A., Carr, T., **Williams, J.**, Panganiban, J., Jones, F., Kimbrough, J., Shih, W., and Kasari, C. (2019), Developmental screening and early intervention in a childcare setting for young children at risk for autism and other developmental delays: A feasibility trial. *Autism Research*, 12(9):1423-1433. doi:10.1002/aur.2160
- Locke, J., **Williams, J.**, Shih, W. and Kasari, C. (2017), Characteristics of socially successful elementary school-aged children with autism. *J Child Psychol Psychiatr*, 58(1):94-102. doi:10.1111/jcp.12636

## Presentations

### 2019 Joint Statistical Meetings

“Maximum Likelihood Estimation of a Truncated Normal Distribution with Censored Data”

Denver, Colorado

### 2019 Western North American Region (WNAR) of the International Biometric Society

“Maximum Likelihood Estimation of a Truncated Normal Distribution with Censored Data”

Portland, Oregon

### 2018 Joint Statistical Meetings

“Propensity Score Methods for Studies with Clustered Data and Continuous Exposure”

Vancouver, BC, Canada

### 2018 Gatlinburg Conference

“Using Clustering to Define ASD Subgroups with Differential Play Outcomes”

San Diego, CA