# Brayan V. Ortiz

#### **Contact:**

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#### **Education:**

University of Washington, Seattle, WA

• Ph.D., Biostatistics, Standard Pathway, 2018

California State University Fullerton, Fullerton, CA

• B.A. Mathematics, (Probability and Statistics Concentration), 2013

## Professional and Research Experience:

Amazon, Modeling and Optimization within Supply Chain Optimization Technologies, Seattle, WA Research Scientist, Modeling & Optimization, July 2018 - present

- Statistical modeling and research
- Reinforcement learning applications in operations
- Integrating optimization-based insight into short and long term planning of transportation network
- Software: Python, FICO Xpress

Centro de Investigaciones de Cancer en Sonora (Sonora Cancer Research Center), Seattle, WA Adjunct Professor of Biostatistics, Biostatistics, 2018 - 2019

• Clinical trial design, statistical analyses, and manuscript preparation.

University of Washington, Department of Biostatistics, Seattle, WA

Research Assistant, Tim Thornton under NIH Statistical Genetics Training Grant, 2013-2014

- Ran principal component analyses to detect population structure such as admixture in genome wide association studies
- Software: R Statistical Software

Research Assistant, Noah Simon, 2014-2015, 2017-2018

- Develop nonparametric penalized regression methodology, which included describing/proving theoretical properties and creating alternating direction method of multipliers solver
- Software: Python (CVXPY, CVXOPT, SciPy, NumPy), C++ (Armadillo), R (Rcpp, RcppArmadillo)

Research Assistant, Jim Hughes, 2015-present

- Develop Bayesian methodology incorporating information from multiple sources to predict pill-taking adherence in HIV prophylaxis clinical trials
- Software: R, OpenBUGS, JAGS, STAN

Teaching Assistant, Biostatistics 310 with Lyndia Brumback, 2017

- Provided supplemental lectures on topics such as data description, study design, screening, estimation hypothesis testing, categorical data analysis, and regression.
- Job duties included lecturing, grading, and preparing lecture slides.

Teaching Assistant, Summer Institute in Statistics for Big Data (SISBID), Visualization of Biomedical Big Data with Dianne Cook and Heike Hoffman

• Provided support during lectures on structured development of static and interactive graphics using ggplot2 in R, especially in the context of exploring big data.

Amazon, Modeling and Optimization within Supply Chain Optimization Technologies, Seattle, WA Research Scientist Intern, Andrew Bruce and Chunyi Wang, June - September 2017

- Collaborated with non-statisticians to define business/financial deliverables followed by collaborations with senior statisticians determining appropriate modeling goals
- Pulled, cleaned, and prepared data to be used in modeling
- Built predictive models to be deployed at a large scale
- Software: R, Python (Pandas, XGBoost), SQL

Yaqui Molecular, Seattle, WA Statistics Consultant, 2015-2016

- Collaborated with clinical immuno-oncologists to determine appropriate statistical methodology for identifying biomarkers associated with cancer outcomes.
- Ran statistical analyses focusing on identifying predictive/prognostic biomarkers in oncology, which includes survival analyses using Cox proportional hazards regression model.
- Based on simulation-based parametric power analyses, provided recommendations for future studies, such as which biomarkers to collect and estimated sample size needed to confidently detect an effect.
- Software: R and Rmarkdown for manuscript preparation

California State University Fullerton, Department of Mathematics, Fullerton, CA

Collaborator with Mori Jomshidian, NIH Funded MARC Program, 2011-2013

- Built logistic regression model, which predicts presence of multiple sclerosis based on performance in cognitive tasking tests.
- Software: R

University of Wisconsin, Madison, Department of Biostatistics, Madison, WI Collaborator with Sushmita Roy, NSF Funded IBS-SRP, 2012

- Ran genome-wide association study to determine association between Alzheimer's cognitive disease and single-nucleotide polymorphisms (SNPs).
- SNP data (> 15 gigabytes) collected from Alzheimer's Disease Neuro-Imaging database.
- Software: PLINK, Python

### Honors, Awards, Scholarships:

- OpsTech Science Fair Grand Prize, "Think Big Award," with Sapphire Manthorpe (co-presenter), 2017
- Trainee, NIH Statistical Genetics Training Grant, University of Washington, 2013-2016
- Minority Access to Research Careers (MARC) Fellowship (NIH), 2011-2013
- CSUF Natural Sciences and Mathematics Symposium Competition Winner, 2012
- SACNAS, Outstanding Poster in Statistics, 2012
- CSUF Special Recognition for Undergraduate Research, 2013
- Joint Mathematics Meeting, San Diego, Outstanding Poster, 2013

### **Publications:**

- Hughes, J., B. Williamson, C. Krakauer, G. Chau, B. Ortiz, J. Wakefield, C. Hendrix, K. Amico, T. Holtz, LG Bekker, & R. Grant, (2020). Combining Information to Estimate Adherence in Studies of Pre-Exposure Prophylaxis for HIV Prevention: Application to HPTN 067. Journal of Statistics in Medicine. *In Submission*
- Ortiz, B. and A. Sinha, (2019). Using Image Transformations to Learn Structure. In progress
- Ortiz, B. and N. Simon, (2017). Mesh-Based Solutions for Nonparametric Penalized Regression. In progress.
- Ortiz, B., M. Jamshidian, and A. Khatoonabadi, (2013). A Statistical Approach to Validate a Cognitive Test for Multiple Sclerosis. Dimensions. Vol. 15. Pp. 105-114
- Ortiz, B., S. Roy, and R. Atlas, (2012). Identification and Characterization of Predictive Genomic Markers in Alzheimer's Disease. IBS-SRP Journal. pp. 121-125

### Conferences

Operations Technology Science Fair (OpsTech Science Fair), Amazon Operations Research. July 2017.
Poster presentation. Amazon Confidential.

•	BayesComp 2018, International Society for Bayesian Based Solutions for Nonparametric Regression."	Analysis.	March 2018.	Poster presentation.	"Mesh