Johns Hopkins Bloomberg School of Public Health
615 N Wolfe St
Office E3032
Baltimore, MD 21205

□ aafisher@hsph.harvard.edu
http://aaronjfisher.github.io

Aaron Fisher

Education

2016 **PhD in Biostatistics**, Johns Hopkins Bloomberg School of Public Health, Baltimore, MD.

Advisors: Vadim Zipunnikov & Brian Caffo

Thesis: Methods for High Dimensional Analysis, Multiple Testing, and Visual Exploration

2010 **BA in Economics**, *University of Rochester*, Rochester, NY. Summa cum laude

Professional Experience

2016-Present

Postdoctoral Research Fellow, Dept of Biostatistics, Harvard T.H. Chan School of Public Health, Boston, MA.

Advisor: Francesca Dominici

2016 Statistical Consultant, Pfizer, Boston, MA.

Analysis of wearable devices and temperature probes in human sleep studies

2010 Intern Analyst, Structured Decisions Corporation, Newton, MA. Background research project for a linear programming application

Academic Papers

Peer-Reviewed Publications

A. J. Fisher, M. Rosenblum (2018). Stochastic Optimization of Adaptive Enrichment Designs for two Subpopulations. (in press) Journal of Biopharmaceutical Statistics. (link)

T. Qian, E. Colantuoni, **A. J. Fisher**, M. Rosenblum, for the Alzheimer's Disease Neuroimaging Initiative (2017). Sensitivity of adaptive enrichment trial designs to accrual rates, time to outcome measurement, and prognostic variables. *Contemporary Clinical Trials Communications*. (<u>link</u>)

Y. Webb-Vargas, S. Chen, **A. J. Fisher**, A. Mejia, Y. Xu, C. Crainiceanu, B. Caffo, M. A. Lindquist (2017). Big Data and Neuroimaging. *Statistics in Biosciences*. (<u>link</u>)

- R. Y. Coley, **A. J. Fisher**, M. Mamawala, H. B. Carter, K. J. Pienta, S. L. Zeger (2017). A Bayesian Hierarchical Model for Prediction of Latent Health States from Multiple Data Sources with Application to Active Surveillance of Prostate Cancer. *Biometrics*. (link)
- M. Rosenblum, T. Qian, Y. Du, H. Qiu, A. J. Fisher (2016). Multiple Testing Procedures for Adaptive Enrichment Designs: Combining Group Sequential and Reallocation Approaches. *Biostatistics*. (<u>link</u>)
- **A. J. Fisher**, B. Caffo, B. Schwartz, V. Zipunnikov (2015). Fast, Exact Bootstrap Principal Component Analysis for p > 1 million. *Journal of the American Statistical Association TM*. (<u>link</u>)
- **A. J. Fisher**, G. B. Anderson, R. Peng, J. Leek (2014). A randomized trial in a massive online open course shows people don't know what a statistically significant relationship looks like, but they can learn. *PeerJ.* (<u>link</u>; 6,428 unique visitors as of September 1, 2015)
- Submitted **A. J. Fisher,** C. Rudin, F. Dominici (2018). Model Class Reliance: Variable Importance Measures for any Machine Learning Model Class, from the "Rashomon" Perspective. (<u>link</u>)
- Technical **A. J. Fisher,** R. Y. Coley, S. L. Zeger (2015). Fast Out-of-Sample Reports Predictions for Bayesian Hierarchical Models of Latent Health States. (<u>link</u>)
 - **A. J. Fisher**, H Jaffee, M Rosenblum (2014). interAdapt An Interactive Tool for Designing and Evaluating Randomized Trials with Adaptive Enrollment Criteria. (<u>link</u>)

Awards and Scholarships

- 2016 Margaret Merrell Award (co-winner with Amanda Mejia): Departmental award recognizing outstanding research by a Biostatistics doctoral student (link).
- 2014 **June B. Culley Award:** Honors outstanding achievement by a Biostatistics student on his or her school-wide oral examination paper (link).
- 2012-2015 **Doctoral Training Grant in Environmental Biostatistics:** Provides funding for at least three years.
- 2006-2010 Undergraduate Awards: Phi Beta Kappa; John Dows Mairs Prize (University of Rochester Economics Dept); Omicron Delta Epsilon International Honor Society for Economics; Theta Chi Long, Walter, Ott Award; Theta Chi Valentine H. Zahn Fund.

Software

bootSVD An R package for implementing fast, exact bootstrap principal component analysis and singular value decompositions for high dimensional data (i.e. > 1 million covariates). Matrices too large for memory can be entered as class ff objects, with contents stored on disk (<u>CRAN link</u>; <u>GitHub link</u>).

ggBrain An R package for beautiful brain image figures with ggplot. This packages allows color to be mapped to both (1) tissue intensities of the template image, and (2) values of a voxel-wise test statistic (GitHub link).

interAdapt An interactive tool for designing and evaluating randomized trials with adaptive enrollment criteria (Shiny App link; CRAN link; Github link).

Computer Skills

Advanced Skills

Basic Skills

git, Python, MATLAB, C, java, D3.js, stata, shell scripting, LATEX

Reviewer

Journal of the American Statistical Association (1); Computational Statistics and Data Science (1); Risk Analysis (2)

Teaching

Co-Instructor

2015 Statistical Reasoning I and II: My role included teaching independently for 13 hours of lectures (MPH Level Course, JHSPH Summer Institute of Epidemiology and Biostatistics)

Guest Lecturer

2013 Essentials of Probability and Statistical Inference I-II (Biostatistics ScM Level, JHSPH)

Lab Lecturer with Content Design

2012-2014 Essentials of Probability and Statistical Inference I-IV: Designed and administered a weekly 1-hour lab lecture. In the second year of this course, we reduced this lab to a 1-hour session every two weeks. (Biostatistics ScM Level, JHSPH)

Lab Lecturer without Content Design

2014-2015 Statistical Methods in Public Health II: Administered approximately 16 hours of lab lecture in each year of the course. (MPH Level, JHSPH)

Educational Presentations

- 2013-2015 JHU Biostatistics Journal Club: I have given talks on high dimensional asymptotics, adaptive clinical trials, and the Bayesian Bootstrap
- 2013-2015 JHU Biostatistics Computing Club: I have given talks on environments in R, and on LaTeX

 General TA Roles
- 2014-2015 Statistical Methods in Public Health I and IV (MPH Level, JHSPH)
 - 2012 Statistical Reasoning I and II, (MPH Level, JHSPH Summer Institute of Epidemiology and Biostatistics)

Conference Presentations

- 2018 "Model Class Reliance: Variable Importance Measures for any Machine Learning Model Class, from the 'Rashomon' Perspective." Harvard Chan Poster Day. *Contributed Poster*.
- 2016 "Optimizing Adaptive Enrichment Designs." JSM, Chicago IL. Contributed Poster.
- 2015 "A Randomized Trial in a Massive Online Open Course Shows People Don't Know What a Statistically Significant Relationship Looks Like, but They Can Learn." JSM, Seattle WA. Contributed Speed Session & Poster.
- 2015 "Fast Exact Bootstrap Principal Component Analysis for p > 1 million." ENAR, Miami, FL. Contributed Talk.
- 2014 "Fast, Exact Bootstrap Principal Component Analysis for p > 1 Million." 4th Annual Hopkins Imaging Conference (<u>link</u>). Baltimore, MD, *Invited Short Talk & Poster*.
- 2014 "Fast Exact Bootstrap Principal Component Analysis for p>1 million: Leveraging Low-Dimensional Structure Across High-Dimensional Bootstrap Samples." JSM, Boston, MA. Contributed Speed Session & Poster.
- 2014 "People Can't See Statistical Significance: A Massive Randomized Trial on the Visual Perception of Relationships." ENAR Spring Meeting, Baltimore, MD, Contributed Talk.

Other Leadership & Service Roles

- 2016 Organizer for HSPH Biostatistics faculty panel on preparing for the job market
- 2015 Volunteer with <u>Thread</u>: Thread is a mentorship and tutoring program that enrolls underperforming high school students who face significant barriers outside of the classroom.

2015	Facilitator at JHU Data Science Hackathon: Assisted a team through
	the process of scraping web data and building a shiny app (3-day event)
2014-2015	JHU Biostatistics Meat Chili Champion
2013-2014	JHU Biostatistics Vegetarian Chili Champion
2012-2013	Co-organizer of JHU Biostatistics Computing Club, with Prasad Patil (speaker schedule link)

Last updated: May 15, 2018