

# Aaron J. Fisher

*Biostatistics & Machine Learning Researcher*

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## Education

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

- Advisors: Vadim Zipunnikov & Brian Caffo
- Awards & Scholarships:
  - Margaret Merrell Award for outstanding research by a Biostatistics doctoral student
  - June B. Culley Award for outstanding achievement on an oral examination paper
  - Doctoral Training Grant in Environmental Biostatistics

**BA in Economics** (2010), *University of Rochester*, Rochester, NY

- Summa cum laude

## Professional experience

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**Principal Statistician** (2019-Present), *Takeda Pharmaceuticals, Statistics and Quantitative Sciences*, Boston, MA

- Analysis of wearable devices in early-stage clinical trials (with Dmitri Volfson)

**Postdoctoral Research Fellow** (2016-2019), *Dept of Biostatistics at the Harvard T.H. Chan School of Public Health*, Boston, MA

- Advisors: Francesca Dominici & Cynthia Rudin

**Statistical Consultant** (2016), *Pfizer*, Boston, MA

- Analysis of wearable devices and temperature probes in human sleep studies

**Intern Analyst** (2010), *Structured Decisions Corporation*, Newton, MA

- Background research project for a linear programming application

## Skills

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**Statistics & Machine Learning:** Causal inference, matrix decompositions, regression in a RKHS, Bayesian regression trees, random forests, neural networks, finite sample bounds, adaptive clinical trials, non-convex quadratic programming, functional data analysis

**Computing:** R package development, git, Python, PyTorch, MATLAB, Stata, shell scripting,  $\LaTeX$

## Summary of selected projects

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**Fast bootstrap PCA for millions of covariates:** In order to quantify uncertainty for the dominant patterns in brain MRI data, with millions of voxels measured for each patient, we developed a fast bootstrap principal component analysis procedure. Our approach reduces computation time from the order of days to the order of minutes, with no loss in accuracy. This work was published in JASA T&M, and my associated R package (bootSVD) has been downloaded over 23,000 times.

**Interpretability for machine learning models:** Black-box, proprietary prediction models can provoke distrust, and produce predictions that are difficult to combine with supple-

mentary information in order to make decisions. In response, we proposed a method for estimating how much unknown proprietary models rely on different covariates, which we applied to the criminal recidivism model COMPAS. The technical aspects of our work combined approaches from finite sample theory, U-statistics, covering numbers, and non-convex quadratic programming. Our paper was published in JMLR, and has been cited over 169 times ([link](#)), including citations to previous arXiv versions.

In more recent work, I have been developing machine learning methods for wearable device data that balance accuracy with interpretability.

**Visual intuition for influence functions used in causal inference:** Influence functions have emerged as a popular framework for combining machine learning with statistical inference, especially within the field of real world (observational) data analysis. Unfortunately, the technical theory underlying influence functions intimidates many researchers away from the subject. In order to make influence functions more approachable, our educational paper, published in *The American Statistician*, builds intuition based on rigorous, visual illustrations. Our hope is that these illustrations can be similarly useful to illustrations of a standard derivative as the “slope at a point.”

**Fast prognostic scores for prostate cancer screening:** As part of a team developing Bayesian risk scores that inform decisions on whether to pursue invasive diagnostic tests, I implemented an importance sampling approach to obtain fast, in-clinic risk updates in response to new patient information.

## [Selected academic papers \(see Google Scholar for full list\)](#)

### [Submitted](#)

**A. J. Fisher** (2020). Treatment effect bias from sample snooping: blinding outcomes is neither necessary nor sufficient. ([link](#).)

### [Peer-Reviewed Publications](#)

**A. J. Fisher**, E. H. Kennedy (2020). Visually Communicating and Teaching Intuition for Influence Functions. *The American Statistician*. ([link](#).)

**A. J. Fisher**, C. Rudin, F. Dominici (2019). All Models are Wrong, but Many are Useful: Learning a Variable’s Importance by Studying an Entire Class of Prediction Models Simultaneously. *The Journal of Machine Learning Research*. ([paper link](#); [169 citations](#) as of July 16, 2020, including citations to previous arXiv versions)

**A. J. Fisher**, M. Rosenblum (2018). Stochastic Optimization of Adaptive Enrichment Designs for two Subpopulations. *Journal of Biopharmaceutical Statistics*. ([link](#).)

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](#).)

**A. J. Fisher**, B. Caffo, B. Schwartz, V. Zipunnikov (2016). Fast, Exact Bootstrap Principal Component Analysis for  $p > 1$  million. *Journal of the American Statistical Association* TM. ([link](#).)

**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*. ([link](#); [10,610 unique visitors](#) as of July 16, 2020.)