

Brian David Williamson

Curriculum Vitae

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Education

- 2019 **Ph.D., Biostatistics**, *University of Washington*.
Committee: Marco Carone (co-chair), Noah Simon (co-chair), Scott Emerson, Peter Gilbert
Dissertation: *A unified approach to model-agnostic variable importance*
- 2017 **M.S., Biostatistics**, *University of Washington*.
- 2014 **B.A., Mathematics**, *Pomona College*.

Professional Experience

Interests: statistical methods for vaccine efficacy trials; statistical methods for biomarker discovery and validation; semiparametric efficiency theory; nonparametric statistics; data science; statistical machine learning

09/2021–
Present **Assistant Investigator**, *Biostatistics Unit, Kaiser Permanente Washington Health Research Institute*.

01/2020–
09/2021 **Post-doctoral research fellow**, *Vaccine and Infectious Disease Division, Fred Hutchinson Cancer Research Center*, Supervisor: Ying Huang.

2015–2019 **Research Assistant**, *Statistical Center for HIV/AIDS Research and Prevention, Fred Hutchinson Cancer Research Center*, Supervisor: James Hughes.

06/2013–
08/2013 **Integrative Cancer Biology Program Research Fellow**, *Stanford University*, Supervisor: Sylvia Plevritis.

Funding

09/2018–
12/2019 *Evaluating predictors of HIV vaccine efficacy: Statistical methods for estimation, testing, and inference*

Principal Investigator: Williamson

NIH award F31AI140836, \$86,459, Impact Score: 10

Honors and Awards

Research Communication and Travel Awards

04/2019 Nonparametrics Section Travel Award, American Statistical Association (ASA)

01/2018 Biometrics Section Travel Award, ASA

06/2017 Most Outstanding Oral Paper Award, Western North American Region (WNAR) of the International Biometric Society

06/2017 Graduate School Fund for Excellence and Innovation Travel Award, University of Washington (UW)

06/2017 Graduate and Professional Student Senate Travel Grant, UW

06/2013 Best Insight Award, UCLA DataFest

Teaching and Service Awards

06/2019 Exceptional Service in Biostatistics Award, UW Department of Biostatistics

09/2018 Excellence in Teaching Award, UW Department of Biostatistics

Academic Honors and Awards

09/2014 Top Scholar Incoming Student Award, UW Department of Biostatistics

05/2014 Distinction in the Senior Exercise, Pomona College

Publications

* denotes joint first-author contribution

14. Huang Y, **Williamson BD**, Moodie Z, Carpp LN, Chambonneau L, DiazGranados CA, and Gilbert PB. Analysis of neutralizing antibodies as a correlate of instantaneous risk of hospitalized dengue in placebo recipients of dengue vaccine efficacy trials. *The Journal of Infectious Diseases*, 2021. doi: 10.1093/infdis/jiab342
13. **Williamson BD**, Hughes JP, and Willis AD. A multi-view model for relative and absolute microbial abundances. *Biometrics*, 2021. doi: 10.1111/biom.13503
12. **Williamson BD**, Magaret CA, Gilbert PB, Nizam S, Simmons C, and Benkeser D. Super LeArner Prediction of NAb Panels (SLAPNAP): a containerized tool for predicting combination monoclonal broadly neutralizing antibody sensitivity. *Bioinformatics*, 2021. doi: 10.1093/bioinformatics/btab398
11. **Williamson BD**, Gilbert PB, Carone M, and Simon N. Rejoinder to “Nonparametric variable importance assessment using machine learning techniques”. *Biometrics*, 2020. doi: 10.1111/biom.13389
10. **Williamson BD**, Gilbert PB, Carone M, and Simon N. Nonparametric variable importance assessment using machine learning techniques (with discussion). *Biometrics*, 2020. doi: 10.1111/biom.13392
9. Duke ER, **Williamson BD**, Borate B, Golob JL, Wychera C, Stevens-Ayers T, Huang M-L, Cossrow N, Wan H, Mast CT, Marks MA, Flowers ME, Jerome KR, Corey L, Gilbert PB, Schiffer JT, and Boeckh MJ. Cytomegalovirus viral load kinetics as surrogate endpoints after allogeneic transplantation. *The Journal of Clinical Investigation*, 2020. doi: 10.1172/JCI133960
8. ***Williamson BD** and *Feng J. Efficient nonparametric statistical inference on population feature importance using Shapley values. In *Proceedings of the 37th International Conference on Machine Learning*, volume 119 of *Proceedings of Machine Learning Research*, pages 10282–10291, 2020. URL <http://proceedings.mlr.press/v119/williamson20a.html>
7. Neidich SD, Fong Y, Li SS, Geraghty DE, **Williamson BD**, Young WC, Goodman D, Seaton KE, Shen X, Sawant S, Zhang L, deCamp AC, Blette BS, Shao M, Yates NL, Feely F, Pyo CW, Ferrari G, Frank I, Karuna ST, Swann E, Mascola J, Graham BS, Hammer SM, Sobieszczyk ME, Corey L, Janes H, McElrath MJ, Gottardo R, Gilbert PB, and Tomaras GD. Antibody Fc effector functions and IgG3 associate with decreased HIV-1 risk. *The Journal of Clinical Investigation*, 2019. doi: 10.1172/JCI126391
6. *Magaret CA, *Benkeser DC, ***Williamson BD**, Borate BR, Carpp LN, Georgiev IS, Setliff I, Dings AS, Simon N, Carone M, Simpkins C, Montefiori D, Alter G, Yu WH, Juraska M, Edlefsen PT, Karuna S, Mgodini NM, Edugupanti S, and Gilbert PB. Prediction of VRC01 neutralization sensitivity by HIV-1 gp160 sequence features. *PLOS Computational Biology*, 2019. doi: 10.1371/journal.pcbi.1006952
5. Hanscom B, Hughes JP, **Williamson BD**, and Donnell D. Adaptive non-inferiority margins under observable non-constancy. *Statistical Methods in Medical Research*, 2018. doi: 10.1177/0962280218801134
4. *Feng J, ***Williamson BD**, Carone M, and Simon N. Nonparametric variable importance using an augmented neural network with multi-task learning. In *Proceedings of the 35th International Conference on Machine Learning*, volume 80 of *Proceedings of Machine Learning Research*, pages 1495–1504, 2018. URL <http://proceedings.mlr.press/v80/feng18a.html>

3. Anchang B, Davis KL, Fienberg H, **Williamson B**, Bendall SC, Karacosta L, Tibshirani R, Nolan GP, and Plevritis SK. DRUG-NEM: optimizing drug combinations using single-cell perturbation response to account for intratumoral heterogeneity. *Proceedings of the National Academy of Sciences*, 115 (18):E4294–E4303, 2018. doi: 10.1073/pnas.1711365115
2. Safren SA, Hughes JP, Mimiaga MJ, Moore AT, Friedman RK, Srithanaviboonchai K, Limbada M, **Williamson BD**, Elharrar V, Cummings V, Magidson JF, Gaydos CA, Celentano D, and Mayer KH for the HPTN063 Study Team. Frequency and predictors of estimated HIV transmissions and bacterial STI acquisition among HIV-positive patients in HIV care across three continents. *Journal of the International AIDS Society*, 19, 2016. doi: 10.7448/IAS.19.1.21096
1. Ritchwood TD, Hughes JP, Jennings L, MacPhail C, **Williamson B**, Selin A, Kahn K, Gómez-Olivé XF, and Pettifor A. Characteristics of age-discordant partnerships associated with HIV risk among young South African women (HPTN 068). *Journal of Acquired Immune Deficiency Syndromes*, 72: 423–429, 2016. doi: 10.1097/QAI.0000000000000988

Preprints

* denotes joint first-author contribution

1. **Williamson BD**, Gilbert PB, Simon NR, and Carone M. A unified approach for inference on algorithm-agnostic variable importance. *arXiv*, 2020. URL <https://arxiv.org/abs/2004.03683>

Statistical Analysis Plans

1. Gilbert PB, Fong Y, Benkeser D, Andriesen J, Borate B, Carone M, Carp LN, Díaz I, Fay MP, Fiore-Gartland A, Hejazi NS, Huang Y, Huang Y, Hyrien O, Janes HE, Juraska M, Li K, Luedtke A, Nason M, Randhawa AK, van der Laan L, **Williamson BD**, Zhang W, and Follman D. USG COVID-19 Response Team / CoVPN Vaccine Efficacy Trial Immune Correlates Statistical Analysis Plan. 2021. doi: 10.6084/m9.figshare.13198595

Software

- `vimp` perform inference on algorithm-agnostic variable importance (available on [CRAN](#))
- `vimpy` perform inference on algorithm-agnostic variable importance in Python (available on [PyPI](#))
- `paramedic` Predicting Absolute and Relative Abundance by Modeling Efficiency to Derive Intervals and Concentrations
- `SLAPNAP` Super Learner Predictions using NAb Panels (available on [DockerHub](#))
- `rigr` regression, inference, and general data analysis tools for [R](#)

Teaching Experience

University of Washington

- Spring 2018 Pre-Doctoral Instructor, Biostatistics 311 – Regression Methods in the Health Sciences (Enrollment: 12; median evaluation: 4.9/5.0; co-taught with Kelsey Grinde)
 - 09/2018, University of Washington School of Public Health Math and R Skills Preparatory Workshop (Enrollment: 175; numerical evaluation not conducted; co-taught with Jessica Williams-Nguyen [2018, 2017] and Gillian Tarr [2016])
 - 09/2017, Williams-Nguyen [2018, 2017] and Gillian Tarr [2016])
 - 09/2016 Williams-Nguyen [2018, 2017] and Gillian Tarr [2016])
- Fall 2017 Lead Teaching Assistant, Biostatistics 511 – Medical Biometry I (Enrollment: 149; numerical evaluation not conducted.)
- Spring 2017 Teaching Assistant, Biostatistics 311 – Regression Methods in the Health Sciences (Enrollment: 22; numerical evaluation not conducted.)
- Winter 2017 Teaching Assistant, Biostatistics 571 – Advanced Regression Methods for Dependent Data (Enrollment: 45; numerical evaluation not conducted)

Pomona College

- Spring 2014 Teaching Assistant, Math 58b – Introduction to Biostatistics
 Fall 2013 Teaching Assistant, Math 58b – Introduction to Statistics

█ Advising and Formal Mentoring

Project Supervision

- 03/2021–08/2021 Drew King, Fred Hutch / Seattle Central College MESA Intern (with Ying Huang)
 02/2021–05/2021 Courtney Simmons, Emory University MS Capstone Project (with David Benkeser)

█ Intramural Service

Fred Hutchinson Cancer Research Center

- 03/2020–09/2021 Hutch United Outreach Committee

Department of Biostatistics, University of Washington

- 02/2019–06/2019 Chair's Task Force on Professionalism; resulted in the [UW Biostatistics Code of Conduct](#)
 09/2016–12/2019 Equity, Diversity, and Inclusion Committee
 09/2016–12/2019 Peer Mentor
 09/2015–09/2018 Educational Policy and Teaching Evaluation Committee

█ Extramural Service

Mentoring

- 2020, 2021 Mentor, Graduate Student Mentorship Initiative, Científico Latino
 06/2015–Present Alumni Mentor, SagePost 47, Pomona College

Manuscript Reviewer

Journal of Machine Learning Research, International Conference on Learning Representations (ICLR; 2022), Journal of the American Statistical Association (Theory & Methods), Neural Information Processing Systems (NeurIPS; 2021), Observational Studies, Data Mining and Knowledge Discovery, Epidemiology, International Journal of Biostatistics

Conferences

- 2020 Program Committee Member, 15th Machine Learning in Computational Biology Meeting
 2019 Program Committee Member, 14th Machine Learning in Computational Biology Meeting
 2017 Session chair, "Quantification, Association Testing, and Integration of the Microbiome", Joint Statistical Meetings

█ Invited Presentations

- 2021 "Inference for Model-Agnostic Variable Importance". Third Annual Hutch United Symposium. Virtual. (*keynote*)

- 2018 “Assessing Variable Importance Nonparametrically using Machine Learning Techniques”. University of Washington Department of Biostatistics Colloquium. Seattle, WA

■ Seminars and Contributed Presentations

- 2021 “Inference for Model-Agnostic Variable Importance”. Kaiser Permanente Washington Health Research Institute Seminar. Virtual.
- 2021 “Statistical Inference and Containerization in Computational Pipelines”. Fred Hutchinson Cancer Research Center Biostatistics Program Seminar. Virtual.
- 2020 “A Unified Approach to Inference on Algorithm-Agnostic Variable Importance”. Vanderbilt University Department of Biostatistics Seminar. Virtual.
- 2020 “Efficient Nonparametric Statistical Inference on Population Feature Importance using Shapley Values”. Thirty-seventh International Conference on Machine Learning. Virtual.
- 2020 “Guiding HIV-1 Antibody Regimen Down-Selection and Prevention Efficacy Trial Design Using Machine Learning”. 27th International Dynamics and Evolution of HIV and Other Human Viruses Meeting. Virtual.
- 2019 “A Unified Approach to Nonparametric Variable Importance Assessment”. Joint Statistical Meetings. Denver, CO (selected for an ASA Nonparametrics Section Travel Award).
- 2018 “Assessing Variable Importance Nonparametrically using Machine Learning Techniques”. Joint Statistical Meetings. Vancouver, BC, Canada (selected for an ASA Biometrics Section Travel Award).
- 2018 “Nonparametric Variable Importance Using an Augmented Neural Network with Multi-Task Learning”. Thirty-fifth International Conference on Machine Learning. Stockholm, Sweden.
- 2017 “Assessing Variable Importance Nonparametrically using Machine Learning Techniques”. WNAR of the International Biometric Society. Santa Fe, NM (selected as the Most Outstanding Oral Paper).
- 2017 “Assessing Variable Importance Nonparametrically using Machine Learning Techniques”. University of Washington Department of Biostatistics Student Seminar. Seattle, WA.
- 2017 “An Introduction to Targeted Learning”. University of Washington Department of Biostatistics Student Seminar. Seattle, WA.
- 2014 “Shrinkage Estimators for High-Dimensional Covariance Matrices”. Pomona College Mathematics Seminar. Claremont, CA.
- 2013 “Automating Cell Gating and Creating a Nested Effects Model to Compare Drug Effects”. Stanford University Center for Cancer Systems Biology Meeting. Stanford, CA.

■ Poster Presentations

- 2018 “Nonparametric Variable Importance using an Augmented Neural Network with Multi-Task Learning”. Thirty-fifth International Conference on Machine Learning. Stockholm, Sweden.
- 2018 “HIV-1 Sequence Predictors of VRC01 Neutralization Sensitivity”. 25th International Dynamics and Evolution of HIV and Other Human Viruses Meeting. Leavenworth, WA.
- 2017 “Assessing Variable Importance Nonparametrically using Machine Learning Techniques”. University of Washington Biostatistics Department Retreat. Seattle, WA.
- 2017 “Assessing Variable Importance Nonparametrically using Machine Learning Techniques”. Joint Statistical Meetings. Baltimore, MD.
- 2015 “Assessing Variable Importance Nonparametrically”. University of Washington Biostatistics Department Retreat. Blaine, WA.

Non-technical, Outreach, and Mentoring Presentations

- 2021 Moderator for “Interdisciplinary Science Panel”. Seattle Central College MESA-LSAMP Program. Virtual.
- 2020 “Statistics in Infectious Disease Research”. Roanoke Valley Governor’s School for Science and Technology Computational Biology Course. Virtual.
- 2020 Panelist for “LSAMP Virtual Research Panel”. Seattle Central College MESA-LSAMP Program. Virtual.
- 2018 “Fellowships, scholarships, and grants”. Biostatistics Student Seminar, University of Washington. Seattle, WA.
- 2017 “Travel grants and conference funding”. University of Washington Department of Biostatistics. Seattle, WA.

Professional Societies

- 08/2014– WNAR of the International Biometric Society
Present
- 03/2013– American Statistical Association
Present
- 05/2014– Sigma Xi: the Scientific Research Honor Society
Present