

# Brian David Williamson

## Curriculum Vitae

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### Biographical Information

Brian D. Williamson, PhD  
Assistant Investigator  
Kaiser Permanente Washington Health Research Institute (KPWHRI)  
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🌐 <https://bdwilliamson.github.io>

### Education

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

### Licensure

Not applicable

### Professional Positions

- 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*.
- 2015–2019 **Research Assistant**, *Statistical Center for HIV/AIDS Research and Prevention, Fred Hutchinson Cancer Research Center*.
- 06/2013–  
08/2013 **Integrative Cancer Biology Program Research Fellow**, *Stanford University*.

### Honors, Awards, and Scholarships

#### 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

### Organizational Service

#### At KPWHRI:

- 11/2021– Present Member, Equity, Inclusion, and Diversity Standing Committee

#### At Fred Hutchinson Cancer Research Center:

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

#### At University of Washington:

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

### External Professional Activities

#### Mentorship roles

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

#### Service in professional associations

- 05/2021– Present Member, Justice, Equity, Diversity, and Inclusion Committee of WNAR

#### 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*

#### Memberships in professional organizations

- 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

## Bibliography

### (a) Refereed research articles

( the symbol ★ denotes joint first-author contribution )

( Published or in press )

- (15) Han S, **Williamson BD**, and Fong Y. Improving random forest predictions in small datasets from two-phase sampling designs. *BMC Biomedical Informatics and Decision Making*, 2021 (*in press*)
- (14) **Williamson BD**, Gilbert PB, Simon NR, and Carone M. A general framework for inference on algorithm-agnostic variable importance. *Journal of the American Statistical Association (Theory & Methods)*, 2021. doi: 10.1080/01621459.2021.2003200 (*in press*)
- (13) 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
- (12) **Williamson BD**, Hughes JP, and Willis AD. A multi-view model for relative and absolute microbial abundances. *Biometrics*, 2021. doi: 10.1111/biom.13503
- (11) **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
- (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, Dingens AS, Simon N, Carone M, Simpkins C, Montefiori D, Alter G, Yu WH, Juraska M, Edlefsen PT, Karuna S, Mgodi 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

( Submitted or under revision )

- (1) Hughes JP, **Williamson BD**, Krakauer C, Chau G, Ortiz B, Wakefield J, Hendrix C, Amico KR, Holtz TH, Bekker LG, and Grant R. Combining information to estimate adherence in studies of pre-exposure prophylaxis for HIV prevention: application to HPTN 067. *Statistics in Medicine*, 2021 (under second round of review)

#### (b) Other refereed scholarly publications

- (1) **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

#### (c) Books and book chapters

#### (d) Other non-refereed scholarly publications

- (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](#)

## Patents and Intellectual Property

Not applicable

## ■■■■■ Funding History

### (a) Current funded projects

- (1) Reducing CNS-active Medications to Prevent Falls and Injuries in Older Adults (U01 CE002967), PI: Phelan, CDC, Co-I, 0.20-0.40 FTE, 09/30/2018–09/29/2023.
- (2) Engaging Staff to Improve COVID-19 Vaccination Rates at Long Term Care Facilities (ENSPIRE), PIs: Hsu and McCracken, PCORI, Co-I, 0.20-0.40 FTE, 8/1/2021–1/31/2025.
- (3) Adaptive Treatment Strategy Methods for Electronic Health Records, PIs: Shortreed and Moodie, NIMH, Co-I, 0.25 FTE, 09/14/2021–09/29/2022.
- (4) Sentinel Advanced Phenotyping, PIs: Carrell and Nelson, FDA, Co-I, 0.15 FTE, 09/14/2021–09/29/2022.

### (b) Completed projects

- (1) Statistical Methods for AIDS Research (R01AI029168), PI: Hughes, NIAID, Research Assistant, 0.5 FTE, 09/2015–09/2018.

## ■■■■■ Conferences and Symposiums

### Conference organization:

1. Program Committee Member, Machine Learning in Computational Biology Meeting, 2019–Present
2. Session chair, “Quantification, Association Testing, and Integration of the Microbiome”, Joint Statistical Meetings, 2017

### Oral presentations: ( the symbol [I] denotes an invited presentation )

16. “Flexible Variable Selection in the Presence of Missing Data”. Joint Statistical Meetings. Virtual, 2021.
15. “Inference for Model-Agnostic Variable Importance”. Third Annual Hutch United Symposium. Virtual, 2021. (*keynote*) [I]
14. “Inference for Model-Agnostic Variable Importance”. Kaiser Permanente Washington Health Research Institute Seminar. Virtual, 2021. [I]
13. “Statistical Inference and Containerization in Computational Pipelines”. Fred Hutchinson Cancer Research Center Biostatistics Program Seminar. Virtual, 2021. [I]
12. “A Unified Approach to Inference on Algorithm-Agnostic Variable Importance”. Vanderbilt University Department of Biostatistics Seminar. Virtual, 2020. [I]
11. “Efficient Nonparametric Statistical Inference on Population Feature Importance using Shapley Values”. Thirty-seventh International Conference on Machine Learning. Virtual, 2020.
10. “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, 2020.
9. “A Unified Approach to Nonparametric Variable Importance Assessment”. Joint Statistical Meetings. Denver, CO, 2019 (selected for an ASA Nonparametrics Section Travel Award).
8. “Assessing Variable Importance Nonparametrically using Machine Learning Techniques”. University of Washington Department of Biostatistics Colloquium. Seattle, WA, 2018. [I]

7. "Assessing Variable Importance Nonparametrically using Machine Learning Techniques". Joint Statistical Meetings. Vancouver, BC, Canada, 2018 (selected for an ASA Biometrics Section Travel Award).
6. "Nonparametric Variable Importance Using an Augmented Neural Network with Multi-Task Learning". Thirty-fifth International Conference on Machine Learning. Stockholm, Sweden, 2018.
5. "Assessing Variable Importance Nonparametrically using Machine Learning Techniques". WNAR of the International Biometric Society. Santa Fe, NM, 2017 (selected as the Most Outstanding Oral Paper).
4. "Assessing Variable Importance Nonparametrically using Machine Learning Techniques". University of Washington Department of Biostatistics Student Seminar. Seattle, WA, 2017.
3. "An Introduction to Targeted Learning". University of Washington Department of Biostatistics Student Seminar. Seattle, WA, 2017.
2. "Shrinkage Estimators for High-Dimensional Covariance Matrices". Pomona College Mathematics Seminar. Claremont, CA, 2014.
1. "Automating Cell Gating and Creating a Nested Effects Model to Compare Drug Effects". Stanford University Center for Cancer Systems Biology Meeting. Stanford, CA, 2013.

#### Poster presentations:

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

#### Non-technical, Outreach, and Mentoring presentations:

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

## ■ Teaching History

### (a) Formal Courses, including Distance Learning

In the University of Washington School of Public Health Math and R Skills Preparatory Workshop

Summer 2016 (co-taught with Gillian Tarr): 175 students

Summer 2017 (co-taught with Jessica Williams-Nguyen): 175 students

Summer 2018 (co-taught with Jessica Williams-Nguyen): 175 students

At the University of Washington

(1) Pre-Doctoral Instructor, Biostatistics 311 – Regression Methods in the Health Sciences

Spring 2018 (co-taught with Kelsey Grinde), enrollment: 12; median evaluation: 4.9/5.0

(b) Other Teaching

(c) Independent Study

## ■■■■■ Advising and Formal Mentoring

(a) Mentored students

(1) Drew King, Fred Hutch / Seattle Central College MESA Intern (with Ying Huang), 03/2021–08/2021

(2) Courtney Simmons, Emory University MS Capstone Project (with David Benkeser), 02/2021–05/2021