

CHARLES J. WOLOCK

Research interests: Nonparametric and semiparametric statistics, survival analysis, missing data, statistical genetics

CONTACT

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Personal webpage	cwoLOCK.github.io
Student webpage	biostat.washington.edu/people/charles-wolock

EDUCATION

University of Washington, Seattle, WA Ph.D., Biostatistics	<i>September 2018 - present</i>
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Harvard University B.A., Organismic and Evolutionary Biology Language citation, Spanish <i>Summa cum laude</i> , Highest Honors Thesis: Exploring the functional diversity of microbial communities within carnivorous pitcher plants Advisors: Anne Pringle, Ph.D. and Naomi Pierce, Ph.D.	<i>September 2011 - May 2015</i>
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RESEARCH EXPERIENCE

Research Assistant University of Washington, Department of Biostatistics Supervisor: Bruce S. Weir, Ph.D.	<i>September 2018 - September 2020</i>
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Research Staff Associate Columbia University, Institute for Genomic Medicine Supervisors: Andrew S. Allen, Ph.D. and David B. Goldstein, Ph.D.	<i>November 2016 - July 2018</i>
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Undergraduate Researcher Harvard University, Department of Organismic and Evolutionary Biology Supervisors: Anne Pringle, Ph.D. and Naomi Pierce, Ph.D.	<i>September 2012 - May 2015</i>
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Harvard University, Department of Earth and Planetary Sciences Supervisor: Erik Sperling, Ph.D.	<i>September 2012 - December 2013</i>
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Summer Scholar Stowers Institute for Medical Research Supervisor: Matthew Gibson, Ph.D.	<i>June 2013 - August 2013</i>
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TEACHING EXPERIENCE

Teaching Assistant University of Washington BIOST 511: Medical Biometry I (Lead TA)	<i>September 2020 - December 2020</i>
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Student Facilitator Harvard University Life Sciences 1a: An Integrated Introduction to the Life Sciences	<i>September 2014 - December 2014</i>
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Tutor

University of Washington

BIOST 523: Statistical Inference for Biometry II

*February 2021 - present***AWARDS, HONORS, FELLOWSHIPS**

NSF Graduate Research Fellowship

*September 2020 - present***University of Washington**

Donovan J. Thompson Award

October 2020

Best combined performance on the PhD Applied and Theory qualifying exams

NIH T32 Statistical Genetics Training Grant

*September 2018 - September 2020***Harvard University**

Phi Beta Kappa

May 2015

Herchel Smith Research Fellowship

June 2014 - August 2014

Microbial Sciences Initiative Research Fellowship

June 2014 - August 2014

John Harvard Scholar

May 2013

National Merit Scholarship

*September 2011***PROFESSIONAL SERVICE**

Manuscript Reviewer

Bayesian Analysis

UNIVERSITY SERVICE

University of Washington, Department of Biostatistics

Student Seminar Coordinator

September 2020 - present

Education Policy and Teaching Evaluation Committee

September 2020 - present

Biostatistics Activities and Events Squad

September 2019 - present

Equity, Diversity, and Inclusion Committee

September 2019 - present

Peer mentor

June 2019 - present

Student-Faculty-Staff Relations Committee

*September 2019 - September 2020***REFEREED JOURNAL PUBLICATIONS**

1. Sperling E.A., **Wolock C.J.**, Morgan A.S., Gill B.C., Kunzmann M., Halverson G.P., Macdonald F.A., Knoll A.H., Johnston D.T. Statistical analysis of iron geochemical data suggests limited late Proterozoic oxygenation. *Nature* 523: 451–454, 2015.
2. Raghavan N.S., Brickman A.M., Andrew H., Manly J.J., Schupf N., Lantigua R., The Alzheimer's Disease Sequencing Project, **Wolock C.J.**, Kamalakaran S., Petrovski S., Tosto G., Vardarajan B.N., Goldstein D.B., Mayeux R. Whole exome sequencing in 20,197 individuals identifies ultra-rare SORL1 loss-of-function variants in late-onset Alzheimer's disease. *Annals of Clinical and Translational Neurology* 5(7): 832-842, 2018.
3. Bittleston L.S., **Wolock C.J.**, Bakhtiar E.Y., Chan X.Y., Chan K.G., Pierce N.E., Pringle A. Convergence between the microcosms of Southeast Asian and North American pitcher plants. *eLife* 7, 2018.
4. Hayeck T.J., Stong N., **Wolock C.J.**, Copeland B., Kamalakaran S., Goldstein D.B., Allen A.S. Improved Pathogenic Variant Localization using a Hierarchical Model of Sub-regional Intolerance. *American Journal of Human Genetics* 104(2): 299-309, 2019.

5. **Wolock C.J.**, Stong N., Ma F., Nagasaki T., Lee W., Tsang S.H., Kamalakaran S., Goldstein D.B., Allikmets R. A case-control collapsing analysis identifies retinal dystrophy genes associated with ophthalmic disease in patients with no pathogenic *ABCA4* mutations. *Genetics in Medicine* 21: 2336-2344, 2019.
6. Gelfman S., Dugger S.A., Moreno C.A.M., Ren Z., **Wolock C.J.**, Shneider N.A., Phatnani H., Cirulli E.T., Lasseigne B.N., Harris T., Maniatis T., Rouleau G.A., Brown R.H., Gitler A.D., Myers R.M., Petrovski S., Allen A.S., Harms M.B., Goldstein D.B. A new approach for rare variation collapsing on functional protein domains implicates specific genic regions in ALS. *Genome Research* 29(5): 809-818, 2019.
7. Cameron-Christie S., **Wolock C.J.**, Groopman E., Petrovski S., Kamalakaran S., Povysil G., Zhang M., Fleckner J., March R.E., Gelfman S., Marasa M., Li Y., Sanna-Cherchi S., Kiryluk K., Allen A.S., Fellström B., Haefliger C., Platt A., Goldstein D.B., Gharavi A. Exome-based rare-variant analyses in chronic kidney disease. *Journal of the American Society of Nephrology* 30(6): 1109-1122, 2019.
8. Ma C.J., **Wolock C.J.**, Stong N., Nagasaki T., Lee W., Goldstein D.B., Allikmets R. Case-control collapsing analysis identifies genes mimicking Stargardt/ABCA4 disease. *Investigative Ophthalmology & Visual Science* 60(9): 2935-2935, 2019.
9. Eade K, Gantner M.L., Hostyk J.A., Nagasaki T., Giles S., Harkins-Perry S., Fallon R., Baldini M., Schepke L., Dorrell M.I., Cai C., Baugh E.H., **Wolock C.J.**, Wallace M., Berlow R.B., Goldstein D.B., Metallo C.M., Friedlander M., Allikmets R. Serine biosynthesis defect due to haploinsufficiency of phosphoglycerate dehydrogenase (PHDGH) causes retinal disease. *Nature Metabolism* 3(3): 366-377, 2021.

SUBMITTED MANUSCRIPTS

1. Wasser S.K., **Wolock C.J.**, Brown J.E., Morris C., Horowitz R., Wong A., Otiende M.Y., Weir B.S. Familial matching of tusks delineates the size and connectivity of transnational criminal organization. Under revision.
2. Bansal A., Heagerty P.J., Inoue L.Y.T., Veenstra D.L., **Wolock C.J.**, Basu A. A Value of Information Framework for Personalizing the Timing of Surveillance Testing. Under revision.

POSTER PRESENTATIONS

1. **Wolock C.J.**, Bittleston L.S., Pierce N.E., Pringle A. Nitrogenase genes in carnivorous plant microbial communities. Microbial Sciences Initiative Research Symposium. Cambridge, MA. September 2014.
2. **Wolock C.J.**, Bittleston L.S., Pierce N.E., Pringle A. Carnivorous pitchers of *Nepenthes* with less acidic fluid house nitrogen-fixing bacteria. Harvard University Organismic and Evolutionary Biology Thesis Symposium. Cambridge, MA. May 2015.
3. **Wolock C.J.**, Kamalakaran S., Goldstein D.B., Allen A.S. A test for balanced coverage across cases and controls as a qualifying criterion in collapsing analysis. Human Genetics in NYC Conference. New York, NY. September 2017.

SKILLS

Computer	Python, R, Bash, SQL, L ^A T _E X
Bioinformatics tools	samtools, bedtools, bcftools, vcftools, Picard tools, BWA, BMap
Languages	English (native), Spanish (proficient)