laron Arbet

UCLA Jonsson Comprehensive Cancer Center

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Research Interests __

- · Machine learning and predictive modeling
- High-dimensional "Big Data"
- Variable selection
- Multi-omics cancer data integration
- Robust nonparametric statistics

Work Experience

Senior Bioinformatician April 2022 - Present

JONSSON COMPREHENSIVE CANCER CENTER, UNIVERSITY OF CALIFORNIA LOS ANGELES, CA

- · Analysis of cancer molecular data: whole genome sequencing, RNA-Seq, DNA methylation, proteomics
- Biomarker identification: among thousands of molecular features, which are most associated with cancer outcomes?
- Daily git/github use for version control and collaboration
- Supervised machine learning for predicting cancer prognosis
- Unsupervised machine learning for deriving cancer subtypes
- · Supervising undegraduate and graduate students on projects
- · Develop open-source R packages for sharing software with broader scientific community
- Create statistical analysis plans and power/sample size estimation for grants

Staff Biostatistician Jan 2021 - April 2022

EDWARDS LIFESCIENCES, IRVINE, CA

- Develop statistical analysis plans and data monitoring for clinical trials
- Power and sample size estimation
- Clinical study reports for federal regulatory agencies
- · Perform analyses with large national database (STS/ACC TVT Registry) including propensity score matching for causal inference

Postdoctoral Research Associate

Aug 2018 - Jan 2021

University of Colorado Anschutz Medical Campus, Department of Biostatistics and Informatics, Aurora, CO

- Provided statistical consulting services
- Published scientific manuscripts
- Developed grants (statistical analysis plan, power/sample size estimation)
- Supervised students and MS/MPH degree biostatisticians
- · Taught BIOS 6601 Intro stats course for 75 MPH students, and created annual 6-week data science machine learning short course
- Develop new statistical methodology

Research Assistant June 2014 - July 2018

University of Minnesota, Division of Biostatistics, Minneapolis, MN

· Developed variable selection methods for high dimensional genetic data and robust methods for estimating the genetic heritability of disease

Statistical Genetics Research Intern

June - Aug 2013

DORDT COLLEGE, DEPARTMENT OF MATHEMATICS, SIOUX CENTER, IA

· Developed methods for aggregating different statistical tests together for greater sensitivity in detecting disease-related genes; resamplingbased bias corrected tests

Math and Statistics Tutor

Sept 2012 - May 2014

WINONA STATE UNIVERSITY, WINONA, MN



University of Minnesota Minneapolis, MN, USA 2014-2018

PHD BIOSTATISTICS

• GPA: 3.7

• GPA: 4.0 | Summa Cum Laude

Peer-reviewed Publications

- Tohi, Y., Sahrmann, J. M., Arbet, J., Kato, T., Lee, L. S., Peacock, M., Ginsburg, K., Pavlovich, C., Carroll, P., Bangma, C. H., Sugimoto, M., & Boutros, P. C. (2025). De-escalation of monitoring in active surveillance for prostate cancer: Results from the GAP3 consortium. *European Urology Oncology*, 8(2), 347–354. https://doi.org/10.1016/j.euo.2024.07.006
- Weiner, A. B., Agrawal, R., Wang, N. K., Sonni, I., Li, E. V., Arbet, J., Zhang, J. J. H., Proudfoot, J. A., Hong, B. H., Davicioni, E., Kane, N., Valle, L. F., Kishan, A. U., Pra, A. D., Ghadjar, P., Sweeney, C. J., Nickols, N. G., Karnes, R. J., Shen, J., Rettig, M. B., Czernin, J., Ross, A. E., Lee Kiang Chua, M., Schaeffer, E. M., Calais, J., Boutros, P. C., & Reiter, R. E. (2024). Molecular hallmarks of prostate-specific membrane antigen in treatment-naïve prostate cancer. European Urology, 86(6), 579–587. https://doi.org/10.1016/j.eururo.2024.09.005
- 3. Lippitt, W., Carlson, N. E., **Arbet, J.**, Fingerlin, T. E., Maier, L. A., & Kechris, K. (2024). Limitations of clustering with PCA and correlated noise. *Journal of Statistical Computation and Simulation*, 94(10), 2291–2319. https://doi.org/10.1080/00949655.2024.2329976
- 4. Ni, K., Rogowitz, E., Farahmand, A. K., Kaizer, L. K., **Arbet, J.**, Cunningham, C. R., Thomas, E. A., & Saxon, D. R. (2024). Weight loss outcomes in a veterans affairs pharmacotherapy-based weight management clinic. *Journal of the Endocrine Society*, 8(5). https://doi.org/10.1210/jendso/bvae042
- 5. Greca, A. L., Grau, L., **Arbet, J.**, Liao, L. M., Sosa, J. A., Haugen, B. R., & Kitahara, C. M. (2023). Anthropometric, dietary, and lifestyle factors and risk of advanced thyroid cancer: The NIH-AARP diet and health cohort study. *Clinical Endocrinology*, 99(6), 586–597. https://doi.org/10.1111/cen.14970
- 6. Gamallat, Y., Choudhry, M., Li, Q., Rokne, J. G., Alhajj, R., Abdelsalam, R., Ghosh, S., **Arbet, J.**, Boutros, P. C., & Bismar, T. A. (2023). Serrate RNA effector molecule (SRRT) is associated with prostate cancer progression and is a predictor of poor prognosis in lethal prostate cancer. *Cancers*, *15*(10), 2867. https://doi.org/10.3390/cancers15102867
- 7. Chan, T. W., Dodson, J. P., **Arbet, J.**, Boutros, P. C., & Xiao, X. (2022). Single-cell analysis in lung adenocarcinoma implicates RNA editing in cancer innate immunity and patient prognosis. *Cancer Research*, 83(3), 374–385. https://doi.org/10.1158/0008-5472.can-22-1062
- 8. Grau, L., **Arbet, J.**, Ostendorf, D. M., Blankenship, J. M., Panter, S. L., Catenacci, V. A., Melanson, E. L., & Creasy, S. A. (2022). Creating an algorithm to identify indices of sleep quantity and quality from a wearable armband in adults. *Sleep Science*, *15*(03), 279–287. https://doi.org/10.5935/1984-0063.20220052
- 9. Creasy, S. A., Ostendorf, D. M., Blankenship, J. M., Grau, L., **Arbet, J.**, Bessesen, D. H., Melanson, E. L., & Catenacci, V. A. (2022). Effect of sleep on weight loss and adherence to diet and physical activity recommendations during an 18-month behavioral weight loss intervention. *International Journal of Obesity*, *46*(8), 1510–1517. https://doi.org/10.1038/s41366-022-01141-z
- 10. Lin, N. W., **Arbet, J.**, Mroz, M. M., Liao, S.-Y., Restrepo, C. I., Mayer, A. S., Li, L., Barkes, B. Q., Schrock, S., Hamzeh, N., Fingerlin, T. E., Carlson, N. E., & Maier, L. A. (2022). Clinical phenotyping in sarcoidosis using cluster analysis. *Respiratory Research*, 23(1). https://doi.org/10.1186/s12931-022-01993-z
- 11. Okamoto, Y., Devoe, S., Seto, N., Minarchick, V., Wilson, T., Rothfuss, H. M., Mohning, M. P., **Arbet, J.**, Kroehl, M., Visser, A., August, J., Thomas, S. M., Charry, L. L., Fleischer, C., Feser, M. L., Frazer? Abel, A. A., Norris, J. M., Cherrington, BrianD., Janssen, W. J., Kaplan, M. J., Deane, K. D., Holers, V. M., & Demoruelle, M. K. (2021). Association of sputum neutrophil extracellular trap subsets with IgA anti–citrullinated protein antibodies in subjects at risk for rheumatoid arthritis. *Arthritis & Rheumatology*, 74(1), 38–48. https://doi.org/10.1002/art.41948
- 12. Wood, C., **Arbet, J.**, Amura, C. R., Nodine, P., Collins, M. R., Orlando, B. S., Mayer, D. C., Stein, D., & Anderson, J. (2021). Multicenter study evaluating nitrous oxide use for labor analgesia at high- and low-altitude institutions. *Anesthesia & Analgesia*, 134(2), 294–302. https://doi.org/10.1213/ane.000000000005712
- 13. Nodine, P. M., **Arbet, J.**, Jenkins, P. A., Rosenthal, L., Carrington, S., Purcell, S. K., Lee, S., & Hoon, S. (2021). Graduate nursing student stressors during the COVID-19 pandemic. *Journal of Professional Nursing*, 37(4), 721–728. https://doi.org/10.1016/j.profnurs.2021.04.008
- 14. **Arbet, J.**, Zhuang, Y., Litkowski, E., Saba, L., & Kechris, K. (2021). Comparing statistical tests for differential network analysis of gene modules. *Frontiers in Genetics*, 12. https://doi.org/10.3389/fgene.2021.630215
- 15. Ostendorf, D. M., Blankenship, J. M., Grau, L., **Arbet, J.**, Mitchell, N. S., Creasy, S. A., Caldwell, A. E., Melanson, E. L., Phelan, S., Bessesen, D. H., & Catenacci, V. A. (2021). Predictors of long-term weight loss trajectories during a

- behavioral weight loss intervention: An exploratory analysis. *Obesity Science & Practice*, 7(5), 569–582. https://doi.org/10.1002/osp4.530
- 16. Reed, S. M., **Arbet, J.**, & Staubli, L. (2021). Clinical nurse specialists in the united states registered with a national provider identifier. *Clinical Nurse Specialist*, 35(3), 119–128. https://doi.org/10.1097/nur.000000000000592
- 17. Rosenthal, L., Lee, S., Jenkins, P., **Arbet, J.**, Carrington, S., Hoon, S., Purcell, S. K., & Nodine, P. (2021). A survey of mental health in graduate nursing students during the COVID-19 pandemic. *Nurse Educator*, 46(4), 215–220. https://doi.org/10.1097/nne.0000000000001013
- 18. Carpenter, C. M., Frank, D. N., Williamson, K., **Arbet, J.**, Wagner, B. D., Kechris, K., & Kroehl, M. E. (2021). tidyMicro: A pipeline for microbiome data analysis and visualization using the tidyverse in r. *BMC Bioinformatics*, 22(1). https://doi.org/10.1186/s12859-021-03967-2
- 19. Ramakrishnan, V. R., **Arbet, J.**, Mace, J. C., Suresh, K., Shintani Smith, S., Soler, Z. M., & Smith, T. L. (2021). Predicting olfactory loss in chronic rhinosinusitis using machine learning. *Chemical Senses*, 46. https://doi.org/10.1093/chemse/bjab042
- 20. Thomas, E. A., Zaman, A., Cornier, M.-A., Catenacci, V. A., Tussey, E. J., Grau, L., **Arbet, J.**, Broussard, J. L., & Rynders, C. A. (2020). Later meal and sleep timing predicts higher percent body fat. *Nutrients*, *13*(1), 73. https://doi.org/10.3390/nu13010073
- 21. Schmanski, A., Roberts, E., Coors, M., Wicks, S. J., **Arbet, J.**, Weber, R., Crooks, K., Barnes, K. C., & Taylor, M. R. G. (2020). Research participant understanding and engagement in an institutional, self-consent biobank model. *Journal of Genetic Counseling*, 30(1), 257–267. https://doi.org/10.1002/jgc4.1316
- 22. **Arbet, J.**, Brokamp, C., Meinzen-Derr, J., Trinkley, K. E., & Spratt, H. M. (2020). Lessons and tips for designing a machine learning study using EHR data. *Journal of Clinical and Translational Science*, 5(1). https://doi.org/10.1017/cts.2020.513
- 23. **Arbet, J.**, McGue, M., & Basu, S. (2020). A robust and unified framework for estimating heritability in twin studies using generalized estimating equations. *Statistics in Medicine*, 39(27), 3897–3913. https://doi.org/10.1002/sim.8564
- 24. Coleman-Minahan, K., Sheeder, J., **Arbet, J.**, & McLemore, M. R. (2020). Interest in medication and aspiration abortion training among colorado nurse practitioners, nurse midwives, and physician assistants. *Women's Health Issues*, 30(3), 167–175. https://doi.org/10.1016/j.whi.2020.02.001
- 25. Gance-Cleveland, B., Linton, A., **Arbet, J.**, Stiller, D., & Sylvain, G. (2020). Predictors of overweight and obesity in childhood cancer survivors. *Journal of Pediatric Oncology Nursing*, 37(3), 154–162. https://doi.org/10.1177/1043454219897102
- 26. James-Allan, L. B., **Arbet, J.**, Teal, S. B., Powell, T. L., & Jansson, T. (2019). Insulin stimulates GLUT4 trafficking to the syncytiotrophoblast basal plasma membrane in the human placenta. *The Journal of Clinical Endocrinology & Metabolism*, 104(9), 4225–4238. https://doi.org/10.1210/jc.2018-02778
- 27. Grinde, K. E., **Arbet, J.**, Green, A., O'Connell, M., Valcarcel, A., Westra, J., & Tintle, N. (2017). Illustrating, quantifying, and correcting for bias in post-hoc analysis of gene-based rare variant tests of association. *Frontiers in Genetics*, 8. https://doi.org/10.3389/fgene.2017.00117
- 28. **Arbet, J.**, McGue, M., Chatterjee, S., & Basu, S. (2017). Resampling-based tests for lasso in genome-wide association studies. *BMC Genetics*, *18*(1). https://doi.org/10.1186/s12863-017-0533-3
- 29. Greco, B., Hainline, A., **Arbet, J.**, Grinde, K., Benitez, A., & Tintle, N. (2015). A general approach for combining diverse rare variant association tests provides improved robustness across a wider range of genetic architectures. *European Journal of Human Genetics*, 24(5), 767–773. https://doi.org/10.1038/ejhg.2015.194

Q Awards

POSTDOCTORAL

Staff appreciation and recognition (STAR)

2023-2024

UCLA JONSSON COMPREHENSIVE CANCER CENTER

Staff appreciation and recognition (STAR)

2022-2023

UCLA JONSSON COMPREHENSIVE CANCER CENTER

■ STUDENT AWARDS

Outstanding Research Assistant Award	2017-2018
University of Minnesota, Division of Biostatistics & Health Data Science	
Interdisciplinary Biostatistics Training Grant in Genetics and Genomics National Institute of General Medical Sciences	2015-2017
COMPETITIONS	
Midwest Undergraduate Data Analysis Competition (MUDAC)	2014
• 3 rd place out of 20 teams from 14 universities	
1st place in the "predictive-modeling class competition" Winona State University Data Mining Course	2014
Midwest Undergraduate Data Analysis Competition (MUDAC)	2013
• 2 nd place out of 15 teams from 10 universities	
Presentations	
■ DEPARTMENTAL	
Robust regression for noisy biological data	
Statistics Working Group, Cancer Data Sciences, University of California, Los Angeles, CA https://jarbet.github.io/presentation-ordinal-regression/ordinal_regression.html	2024
Linear regression diagnostics	
BOUTROS LAB, CANCER DATA SCIENCES, UNIVERSITY OF CALIFORNIA, LOS ANGELES, CA • https://jarbet.github.io/presentation-regression-diagnostics/regression_diagnostics.html	2024
Causal inference with observational data using propensity score matching	
Statistics Working Group, Cancer Data Sciences, University of California, Los Angeles, CA https://jarbet.github.io/presentation-causal-inference-matching/causal_inference_matching.html	2024
Automatic feature selection and engineering with MARS	
STATISTICS WORKING GROUP, CANCER DATA SCIENCES, UNIVERSITY OF CALIFORNIA, LOS ANGELES, CA	2023
• https://jarbet.github.io/presentation-mars.github.io/presentation-mars.html	
Introduction to Bayesian statistics Boutros Lab, Cancer Data Sciences, University of California, Los Angeles, CA	2022
• https://jarbet.github.io/presentations/2022-08-25_Bayesian_stats.pdf	2022
Multiomics cancer data analysis	
Boutros Lab, Cancer Data Sciences, University of California, Los Angeles, CA	2021
• https://jarbet.github.io/presentations/2021-11-21_multiomics_cancer.pdf	
Interpretable machine learning	
Center for Innovative Design and Analysis, University of Colorado, Denver, CO	2021
 https://jarbet.github.io/presentations/2020-01-21_interpretable_machine_learning.pdf 	
Differential Network Analysis of Gene Modules	0004
Systems Genetics and Bioinformatics Research Lab, University of Colorado, Denver, CO	2021
Frequentist versus Bayesian Approaches to Statistics: A Debate Department of Biostatistics and Informatics, University of Colorado, Denver, CO	2018
Robust Estimation of Genetic and Environmental Variance Components in Twin Studies	2010
University of Minnesota Biostatistics Student Seminar Series, Minneapolis, MN	2017
A General framework for Association Tests with Multivariate Traits in Large-Scale	
Genomics Studies	
University of Minnesota Statistical Genetics Journal Club, Minneapolis, MN	2016

Methods of Inference for Penalized Regression in High-Dimensional Genetic Association Studies

School of Public Health Research Day, University of Minnesota, Minneapolis, MN 2016

Penalized Regression for High-Dimensional Genetic Association Testing

University of Minnesota Statistical Genetics Journal Club, Minneapolis, MN

CONFERENCES

Methylome-Genome Interactions Define Disease Aggression in Localized Prostate Cancer

RECOMB/ISCB Conference on Regulatory and Systems Genomics, Los Angeles, CA

2024

Poster

Triplet matching: propensity score matching with 3 groups

Talk

14TH ANNUAL FDA/ADVAMED MEDICAL DEVICE STATISTICAL ISSUES CONFERENCE (VIRTUAL)

2022

 $•\ https://jarbet.github.io/presentations/20220422_triplet_matching.pdf$

Statistical Considerations and Methods to Utilize Real World Evidence in Medical Device Evaluation

Panel Co-organizer

14TH ANNUAL FDA/ADVAMED MEDICAL DEVICE STATISTICAL ISSUES CONFERENCE (VIRTUAL)

2022

Robust Estimation of Genetic and Environmental Variance Components in Twin Studies using Generalized Estimating Equations

Poster

JOINT STATISTICAL MEETINGS (JSM), BALTIMORE, MD

2017

Methods of Inference for Penalized Regression in High-Dimensional Genetic Association Studies

Poster

EASTERN NORTH AMERICAN REGION INTERNATIONAL BIOMETRIC SOCIETY (ENAR), AUSTIN, TX

2016

What now? Post-Hoc Approaches for Gene-based Rare-Variant Tests of Association

Poster

AMERICAN SOCIETY OF HUMAN GENETICS CONFERENCE (ASHG), BOSTON, MA

2013

General Approaches for Combining Rare Variant Association Tests Provide Improved Power Across a Wide Range of Genetic Architectures

Poster

AMERICAN SOCIETY OF HUMAN GENETICS CONFERENCE (ASHG), BOSTON, MA

2013

♣ Teaching Experience _



BIOS 6601 – Applied Biostatistics I

Fall 2019

University of Colorado Anschutz Medical Campus

- Duty: Sole Instructor, completely redesigned all lectures, homework, quizzes, and exams
- **Description**: Applied biostatistical methods including descriptive and statistical inference; odds ratio and relative risk, probability theory, parameter estimation, tests for comparing statistics of two or more groups, correlation, linear regression, logistic regression and survival analysis.
- Credits: 3
- Audience: 75 Masters of Public Health (MPH) students

■ SHORT COURSES

Fundamentals of Data Science Literacy

Summer 2019, Fall 2020

University of Colorado Anschutz Medical Campus

- Duty: lead organizer, developed and taught 3 machine learning lectures
- Description: 6-week short course (2 hours/week) on data science with emphasis on how machine learning and data mining are used in biomedical research. Topics included: 1) Overview of Data Science; 2) Introduction of Machine Learning in Biomedical Research; 3) Clustering and Pattern Finding (unsupervised Machine Learning); 4) Powerful Predictive Models and Variable Selection (supervised Machine Learning); 5) Fundamentals of Data Visualization; 6): Reproducible Research and Team Science
- Audience: 20 medical researchers, mostly non-statisticians

Introduction to Statistics Summer 2019, 2020

University of Colorado Anschutz Medical Campus

- · Duty: Instructor, developed and taught lectures on "Introduction to statistics", study design, and survey design
- Description: Introductory statistics day-long short course for UCHealth nurses
- Audience: 15 UCHealth Nurses

■ GUEST LECTURES

IPHY 5900 - Data Literacy in Biomedical Research

University of Colorado Boulder

- Primary Instructor: Dr. Celine Marie Claire D. Vetter, Integrative Physiology
- Lecture: Machine Learning in Biomedical Research

BIOS 6611 - Biostatistical Methods I

Fall 2018

Nov 2019, Oct 2020

University of Colorado School of Public Health

- Primary Instructor: Dr. Alex Kaizer
- Lecture: Linear Algebra Review

Mentoring and supervising

■ University of California Los Angeles

Supervising students for data analysis projects

- Yuxi Song
- · Sattwik Banerjee
- · Adriana Wiggins

■ University of Colorado, Department of Biostatistics and Informatics

Supervising MPH/MS Statisticians

- Laura Grau (MPH), Nov 2018 Jan 2021
 - I supervised Laura on several projects which led to publications.

Supervising students for data analysis projects

- Yaxu Zhuang (PhD), Jan 2019 Aug 2020
- Rachel Weber (MS), Dec 2018 April 2019

Thesis or Dissertation Committee Member (graduating year)

- Charlie Carpenter, Biostatistics MS thesis committee, Biostatistics (2020)
- Rachel Blumhagen, Biostatistics PhD dissertation committee (2021)

Professional Service

GRANT AND PROTOCOL REVIEWS

Scientific Advisory & Review Committee (SARC), Colorado Clinical and Translational Sciences Institute, University of Colorado – April 2019 to Jan 2021

JOURNAL REVIEWER

- Biometrics
- Genetic Epidemiology
- PeerJ

■ COMMITTEES/OTHER

Department of Biostatistics and Informatics, University of Colorado

- Student Admissions Committee, Fall 2018 Spring 2019
- Volunteer Judge for CIDA Hackathon (24 hour data analysis competition), Jan. 2020

Biostatistics, Epidemiology, & Research Design Core of the Colorado Clinical and Translational Sciences Institute, University of Colorado

• Education Committee, Fall 2018 – Jan 2020

☐ Software Skills_____

PROFICIENT

R

- Over 14 years of experience with R for data analysis
- I enjoy **developing R packages** to easily share documented code with colleagues and the broader research community. Examples:
 - https://uclahs-cds.github.io/package-PrCaMethy/
 - https://github.com/arbet003/discoMod
- Parallel programming in R using multiple cpus

git/github

- Version control
- Used daily at UCLA
- · Collaborating with others on projects
- Review colleagues' code
- Assign issues to team members

Quarto and Rmarkdown

- Reproducible research, manuscripts and reports
- Reproducible HTML presentations. Examples: https://jarbet.github.io/presentations
- Websites: https://jarbet.github.io/
- Resume: this resume was made using Rmarkdown with the vitae R package

Bayesian statistical programming

• Proficient with BUGS/JAGS; basic skills with STAN

■ BASIC SKILLS

- Linux command line and bash scripts
 - Daily use of a high-performance computing cluster at UCLA
- Databricks
- · Microsoft Office
- SAS