Jordan Grey Bryan

Email: jbry@unc.edu http://j-g-b.github.io/ Mobile: +1-310-720-2236

Work Experience

UNC Department of Biostatistics

Postdoctoral trainee

Chapel Hill, NC May 2023 - present

Cambridge, MA

Sep 2015 - Jul 2018

Broad Institute of MIT and Harvard

Associate Computational Biologist I - II

- o Cancer Data Science: Used statistical learning methods to find links between cancer genomics and treatment response in CRISPR knockout screens. Formulated modified mixed effects model and built optimizer to correct for variable sensitivity to cutting-induced DNA damage. Maintained team R package.
- PRISM: Lead quality control and interpretation of data from the PRISM high-throughput screening technology. Wrote software to automate tailored analyses for scientists who submitted compounds to quarterly screens. Presented hypotheses regarding compound mechanism of action to collaborators in industry and academia.
- Vice Chair, Project Brooklyn: Recruited to task-force charged with reforming laboratory and office space management. Partnered with Brightspot Strategy to design prototypes of new library and co-working spaces. Reported findings and made recommendations to Executive Leadership Team at Broad.

EDUCATION

Duke University

PhD, Statistics; Certificate in Slavic, Eurasian, and East European Studies

Durham, NC

Sep 2018 - May 2023

Stanford University

BS, Mathematics with Honors in the Arts

Stanford, CA

Sep 2011 - Jun 2015

Publications

- Bryan, J.G., Hoff P.D., Osburn C.L. (2023) "Routine estimation of dissolved organic matter sources using fluorescence data and linear least squares." ACS ES&T Water.
- de Matos Simoes, R., Shirasaki, R. [& 39 others including Bryan, J.G.]. (2023) "Genome-scale functional genomics identify genes preferentially essential for multiple myeloma cells compared to other neoplasias." Nature Cancer.
- Bryan, J.G., Niles-Weed, J., Hoff P.D. (2022) "The multirank likelihood for semiparametric canonical correlation analysis." Submitted to the Journal of Multivariate Analysis, https://arxiv.org/abs/2112.07465
- Bryan, J.G., Hoff, P.D. (2021) "Smaller p-values in genomics studies using distilled auxiliary information." Biostatistics.
- Sheffer, M., Lowry, E. [& 29 others including Bryan, J.G.]. (2021) "Genome-scale screens identify factors regulating tumor cell responses to natural killer cells." Nature Genetics.
- Bryan, J.G., Mandan, A. [& 8 others]. (2021) "Likelihood ratio statistics for gene set enrichment in Alzheimer's disease pathways." Alzheimer's & Dementia.
- Dhimolea, E., Simoes, R.D.M. [& 29 others including Bryan, J.G.]. (2021) "An Embryonic Diapause-like Adaptation with Suppressed Myc Activity Enables Tumor Treatment Persistence." Cancer Cell.
- Wu, J., Bryan, J. G. [& 10 others]. (2020) "Opportunities and Challenges for Analyzing Cancer Data at the Inter- and Intra-Institutional Levels." JCO Precision Oncology.
- Corsello, S.M., Nagari, R.T. [& 35 others including Bryan, J.G.]. (2020) "Discovering the anticancer potential of non-oncology drugs by systematic viability profiling." Nature Cancer.
- McFarland, J.M., Ho, Z.V. [& 12 others including Bryan, J.G.]. (2018) "Improved Estimation of Cancer Dependencies from Large-Scale RNAi Screens Using Model-Based Normalization and Data Integration." Nature Communications.
- Gray, S.W., Gagan, J. [& 12 others including Bryan, J.G.]. (2018) "Interactive or static reports to guide clinical interpretation of cancer genomics." Journal of the American Medical Informatics Association.
- Meyers, R.M., Bryan, J.G. [& 29 others]. (2017) "Computational correction of copy-number effect improves specificity of CRISPR-Cas9 essentiality screens in cancer cells." Nature Genetics.
- Mackey, L., Bryan, J.G. & Mo, M.Y. (2015) "Weighted Classification Cascades for Optimizing Discovery Significance in the HiggsML Challenge." Journal of Machine Learning Research, Workshop and Conference Proceedings.

Additional Research Experience

Duke Molecular Physiology Institute (DMPI)

Durham, NC

Statistician in Residence

Jun 2022 - Jun 2023

• Office hours: Held weekly office hours for scientists at DMPI. Assisted with data processing in R and advised practitioners on use of statistical methodologies in research.

American Statistical Association

Alexandria, VA

Research Intern

May 2021 - Aug 2021

• State of the federal workplace: Gathered payroll and employee survey data from federal and third-party websites. Ran regression models and applied dimension reduction techniques to interpret the relationship between payroll statistics and employee satisfaction.

Stanford University Department of Statistics

Stanford, CA

Research Assistant

Jun 2014 - Aug 2014

• **Kaggle prediction challenge**: Competed in the HiggsML challenge with the objective of classifying particle decay based on data simulated from the Large Hadron Collider at CERN.

Teaching

- Head teaching assistant and lab instructor: STA 101: Data Analysis and Statistical Inference (2018); STA 360: Bayesian Inference and Modern Statistical Methods (2019-2020); STA 199: Introduction to Data Science (2022)
- Teaching assistant and lab instructor: STA 325: Machine Learning and Data Mining (2020); STA 521: Predictive Modeling and Statistical Learning (2021)

ACADEMIC SERVICE

- Statistics translator: Joined collaborative effort organized by The Society for Imprecise Probabilities: Theories and Applications to translate V.P. Kuznetsov's *Interval Statistical Models* from Russian to English (2022-present).
- GCC Co-Chair: Served on Duke Statistical Science Graduate Consultative Committee (2019-2020, 2022-2023).
- Journal review: Annals of Applied Statistics (AOAS), Alzheimer's & Dementia, Statistical Papers, Statistics and Its Interface.

AWARDS

Danish Data Science Academy, Visit Grant(2023)	Stanford Wrestling, Coaches Award (2015)
ISBA, Graduate Student Travel Award (2022)	Rhodes Scholarship Finalist, District 16 (2014)
BNP Networking Event, Travel Grant (2022)	Stanford Wrestling, Todd Surmon Award (2012)
Duke University, GSTEG Summer Grant (2021)	Stanford Wrestling, Coaches Award (2012)