

Jacob Fiksel

CONTACT INFORMATION	1019 N. Calvert St Baltimore, MD 21202	608-345-8988 jfiksel@gmail.com
RESEARCH INTERESTS	Applied Bayesian statistical modeling, global public health, cancer genomics	
EDUCATION	Johns Hopkins Bloomberg School of Public Health , Baltimore, MD Ph.D., Biostatistics, <i>Expected</i> : Spring 2020 <ul style="list-style-type: none">• Thesis Topic: Bayesian Quantification Learning with Applications to Mortality Surveillance• Advisor: Abhirup Datta Ph.D Pomona College , Claremont, CA B.S., Mathematics with a Statistics focus, May 2015 <ul style="list-style-type: none">• <i>Cum Laude</i>• GPA 3.85/4.00• Distinction in the Senior Exercise	
RESEARCH EXPERIENCE	Johns Hopkins Bloomberg School of Public Health <ul style="list-style-type: none">• Developed a novel Bayesian framework using generalized Gibbs updates for robust quantification• As part of the Countrywide Mortality Surveillance for Action (COMSA) in Mozambique team, built the calibratedVA R package for application of the Bayesian quantification framework to estimation of the leading causes of neonatal and child death in Mozambique.• Assisted in the development and implementation of statistical methods to detect chromosomal abnormalities in circulating cell-free DNA collected from cancer patients. This work resulted in a Nature publication• In collaboration with a clinical radiation oncologist, constructed a Random Survival Forest that uses clinical data from patients with bone metastases to make personalized estimates of survival, allowing for more informed decision making in the choice of palliative radiation treatment regimen• Developed widely-used teaching guides for the use of GitHub Classroom in teaching undergraduate statistics.	
REFEREED JOURNAL PUBLICATIONS	<ol style="list-style-type: none">1. Jillian Phallen, Alessandro Leal, Brian D Woodward, Patrick M Forde, Jarushka Naidoo, Kristen A Marrone, Julie R Brahmer, Jacob Fiksel, Jamie E Medina, Stephen Cristiano, et al. Early noninvasive detection of response to targeted therapy in non-small cell lung cancer. <i>Cancer research</i>, 79(6):1204–1213, 20192. Stephen Cristiano*, Alessandro Leal*, Jillian Phallen*, Jacob Fiksel*, Vilmos Adleff, Daniel C Bruhm, Sarah Østrup Jensen, Jamie E Medina, Carolyn Hruban, James R White, et al. Genome-wide cell-free dna fragmentation in patients with cancer. <i>Nature</i>, page 1, 20193. Jacob Fiksel, Leah R Jager, Johanna S Hardin, and Margaret A Taub. Using github classroom to teach statistics. <i>Journal of Statistics Education</i>, pages 1–10, 20194. Sara R Alcorn, Christen Elledge, Jean L Wright, Thomas J Smith, Todd R McNutt, Jacob Fiksel, Scott Zeger, and Theodore L DeWeese. Frequency of complicated symptomatic bone metastasis over a breadth of operational definitions. <i>International Journal of Radiation Oncology · Biology · Physics</i>, 2019	

* Co-first author

	<p>5. Abhirup Datta, Jacob Fiksel, Agbessi Amouzou, and Scott Zeger. Regularized Bayesian transfer learning for population level etiological distributions. <i>arXiv e-prints</i>, page arXiv:1810.10572, Oct 2018. Accepted in <i>Biostatistics</i></p>
PAPERS IN REVIEW	<p>1. Jacob Fiksel, Abhirup Datta, Agbessi Amouzou, and Scott Zeger. Generalized Bayesian Quantification Learning. <i>arXiv e-prints</i>, page arXiv:2001.05360, Jan 2020</p> <p>2. Sara R. Alcorn, Jacob Fiksel, Jean L. Wright, Christen R. Elledge, Thomas J. Smith, Powell Perng, Sarah Saleemi, Todd McNutt, Theodore L. DeWeese, and Scott L. Zeger. Developing an Improved Statistical Approach for Survival Estimation in Bone Metastases Management: The Bone Metastases Ensemble Trees for Survival (BMETS) Model. Under review at <i>International Journal of Radiation Oncology · Biology · Physics</i></p>
PAPERS IN PREPARATION	<p>1. Jacob Fiksel, Abhirup Datta, and Scott Zeger. A Transformation-free Linear Regression for Compositional Outcomes and Predictors</p>
COMPUTING KNOWLEDGE	<ul style="list-style-type: none"> • R coding and package development • Stan, JAGS, and custom MCMC samplers • Development of machine learning models • Next generation sequencing analysis with command line tools and R • Reproducible computing with Git, GitHub, and GitHub Classroom • Simulation studies using a high performance computing exchange (JHPCE)
PRESENTATIONS	<p>Johns Hopkins Research on Aging Showcase Poster Competition April 2018</p> <ul style="list-style-type: none"> • <i>Optimized Survival Evaluation to Guide Bone Metastases Management: Developing an Improved Statistical Approach</i>. 1st place in the graduate student poster competition
TEACHING EXPERIENCE	<p>Lead Teaching Assistant</p> <ul style="list-style-type: none"> • Fall 2018, 2019: Statistical Methods in Public Health I-II <ul style="list-style-type: none"> – Lead weekly labs 2-3 times per week for 10-40 graduate students <p>Teaching Assistant</p> <ul style="list-style-type: none"> • Spring 2018: Analysis of Longitudinal Data and Multilevel Statistical Models in Public Health • Spring 2017: Statistical Methods in Public Health III-IV • Fall 2016, 2018: Biostatistics for Public Health <ul style="list-style-type: none"> – Lead weekly labs once per week for 25 undergraduate students
SERVICE	<p>JHSPH Biostatistics Computing Club Co-President September 2016-June 2017</p> <ul style="list-style-type: none"> • Organized bi-weekly student presentations on computing related topics for the Biostatistics department <p>Recruiting Committee, Division of Biostatistics December 2016 – Present</p> <ul style="list-style-type: none"> • Assist with planning of annual Division of Biostatistics Open House and Admitted Student Visit Days • Meet with prospective and admitted students

REFERENCES

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