NICK STRAYER

I have made visualizations viewed by hundreds of thousands of people, sped up query times for 25 terabytes of data by an average of 4,800 times, and built packages for R that let you do magic.

EDUCATION PhD. Candidate, Biostatistics 2020 ♥ Nashville, TN Vanderbilt University 2015 · Focused on network models & interactive visualization platforms for electronic health records data University Graduate Fellow B.S., Mathematics, Statistics (minor C.S.) 2015 Burlington, VT University of Vermont 2011 Thesis: An agent based model of Diel Vertical Migration patterns of Mysis diluviana **I** INDUSTRY EXPERIENCE Software Engineer Current **Q** Remote RStudio 2020 • Helping make programming web applications with R easier and more beautiful on the Shiny team Data Journalist - Graphics Department 2016 • New York, New York New York Times 2016 · Reporter with the graphics desk covering topics in science, politics, and sport. • Work primarily done in R, Javascript, and Adobe Illustrator. **Engineering Intern - User Experience** 2015 Burlington, VT Dealer.com 2015 · Built internal tool to help analyze and visualize user interaction with back-end products. 2015 **Data Science Intern** Burlington, VT Dealer.com 2015 · Worked with the product analytics team to help parse and visualize large stores of data to drive business decisions.



View this CV online with links at nickstrayer.me/cv/

CONTACT

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- in linkedin.com/in/nickstrayer

LANGUAGE SKILLS



Made with the R package **page**down.

The source code is available on github.com/nstrayer/cv.

Last updated on 2020-11-04.

2015 2014	•	Data Artist In Residence Conduce Carpinteria, CA • Envisioned, prototyped and implemented visualization framework in the course of one month. • Constructed training protocol for bringing third parties up to speed with new protocol.
2014 2014	•	Software Engineering Intern Conduce Queries Carpinteria, CA • Incorporated d3.js to the company's main software platform.
	₽	RESEARCH EXPERIENCE
Current 2015	•	Graduate Research Assistant TBILab (Yaomin Xu's Lab) ♥Vanderbilt University • Primarily working with large EHR and Biobank datasets. • Developing network-based methods to investigate and visualize clinical- ly relevant patterns in data.
2018 2017	•	Data Science Researcher ♥ Johns Hopkins University Data Science Lab ♥ Johns Hopkins University • Building R Shiny applications in the contexts of wearables and statistics education. • Work primarily done in R Shiny and Javascript (node and d3js).
2015 2013	•	 Undergraduate Researcher Rubenstein Ecosystems Science Laboratory Analyzed and visualized data for CATOS fish tracking project. Head of data mining project to establish temporal trends in population densities of Mysis diluviana (Mysis). Ran project to mathematically model the migration patterns of Mysis (honors thesis project.)
2015 2015	•	Human Computer Interaction Researcher LabInTheWild (Reineke Lab) • Led development and implementation of interactive data visualizations to help users compare themselves to other demographics.
2014 2013	•	Undergraduate Researcher Bentil Laboratory ♥ University of Vermont • Developed mathematical model to predict the transport of sulfur through the environment with applications in waste cleanup.

2013 | 2012

Research Assistant

Adair Laboratory

Ouniversity of Vermont

• Independently analyzed and constructed statistical models for large data sets pertaining to carbon decomposition rates.

TEACHING EXPERIENCE

2020	•	Javascript for Shiny Users
		RStudio::conf 2020
		 Served as TA for two day workshop on how to leverage Javascript in Shiny applications
		 Lectured on using R2D3 package to build interactive visualizations.
2019	•	Data Visualization Best Practices
l 2019		DataCamp
		Designed from bottom up course to teach best practices for scientific visualizations. Uses D and gaplet?
		 In top 10% on platform by popularity.
2019	•	Improving your visualization in Python
 2019		DataCamp
2015		 Designed from bottom up course to teach advanced methods for en- hancing visualization.
		 Uses python, matplotlib, and seaborn.
2018	•	Advanced Statistical Learning and Inference
 2017		Vanderbilt Biostatistics Department • Nashville, TN
2017		• TA and lectured
		 Topics covered from penalized regression to boosted trees and neural networks
		• Highest level course offered in department
2018	•	Advanced Statistical Computing
 2018		Vanderbilt Biostatistics Department Q Nashville, TN
		• TA and lectured
		 Covered modern statistical computing algorithms 4th year PhD level class
2017	•	Statistical Computing in R
 2017		Vanderbilt Biostatistics Department Q Nashville, TN
		• TA and lectured
	I	 Covered introduction to R language for statistics applications Graduate level class

I am passionate about education. I believe that no topic is too complex if the teacher is empathetic and willing to think about new methods of approaching task.

	SELECTED DATA SCIENCE WRITING	
2019	• Using AWK and R to Parse 25tb LiveFreeOrDichotomize.com	
	 Story of parsing large amounts of genomics data. Provided advice for dealing with data much larger than disk. Reached top of HackerNews. 	I regularly blog about data sci- ence and visualization on my blog LiveFreeOrDichotomize
2018	• Classifying physical activity from smartphone data RStudio Tensorflow Blog	
	 Walk through of training a convolutional neural network to achieve stat of the art recognition of activities from accelerometer data. Contracted article. 	e
2018	• The United States of Seasons LiveFreeOrDichotomize.com	
	 GIS analysis of weather data to find the most 'seasonal' locations in Unit ed States Used Bayesian regression methods for smoothing sparse geospatial data. 	t-
2017	• A year as told by fitbit	
	 Analyzing a full years worth of second-level heart rate data from wear- able device. Demonstrated visualization-based inference for large data. 	
2017	• MCMC and the case of the spilled seeds	
	LiveFreeOrDichotomize.com	
	 Full Bayesian MCMC sampler running in your browser. Coded from scratch in vanilla Javascript. 	
2017	• The Traveling Metallurgist LiveFreeOrDichotomize.com	
	 Pure javascript implementation of traveling salesman solution using simulated annealing. 	
	Allows reader to customize the number and location of cities to attempt to trick the algorithm.	t
	SELECTED PRESS (ABOUT)	
2017 	• Great paper? Swipe right on the new 'Tinder for preprints' app	
2017	• Story of the app Papr made with Jeff Leek and Lucy D'Agostino	
	McGowan.	

2017 2017	 Swipe right for science: Papr app is 'Tinder for preprints' Nature News Second press article for app Papr.
2016 2016	 The Deeper Story in the Data University of Vermont Quarterly Story on my path post graduation and the power of narrative.



■ SELECTED PRESS (BY)

2016	•	The Great Student Migration
2016		The New York Times
2010		• Most shared and discussed article from the New York Times for August 2016.
2016	•	Wildfires are Getting Worse, The New York Times
		The New York Times
2010		 GIS analysis and modeling of fire patterns and trends Data in collaboration with NASA and USGS
2016		Who's Speaking at the Democratic National Convention?
		The New York Times
2016		 Data scraped from CSPAN records to figure out who talked and past conventions.
2016	•	Who's Speaking at the Republican National Convention?
 2016		The New York Times
2010		 Used same data scraping techniques as Who's Speaking at the Democ- ratic National Convention?
2016	•	A Trail of Terror in Nice, Block by Block
		The New York Times
2018		 Led research effort to put together story of 2016 terrorist attack in Nice, France in less than 12 hours.
		• Work won Silver medal at Malofiej 2017, and gold at Society of News and Design.

		SELECTED PUBLICATIONS, POSTERS, AND TALKS
2020	•	Building a software package in tandem with machine learning methods research can result in both more rigorous code and more rigorous research ENAR 2020 • Invited talk in Human Data Interaction section. • How and why building an R package can benefit methodological
2020	•	 research Stochastic Block Modeling in R, Statistically rigorous clustering with rigorous code RStudio::conf 2020 Invited talk about new sbmR package. Focus on how software development and methodological research can improve both benefit when done in tandem.
2020	•	 PheWAS-ME: A web-app for interactive exploration of multimorbidity patterns in PheWAS Bioinformatics Manuscript detailing application for the exploration of multimorbidity patterns in PheWAS analyses See landing page for more information.
2019 2019	•	Charge Reductions Associated with Shortening Time to Recovery in Septic Shock Chest • Authored with Wesley H. Self, MD MPH; Dandan Liu, PhD; Stephan Russ, MD, MPH; Michael J. Ward, MD, PhD, MBA; Nathan I. Shapiro, MD, MPH; Todd W. Rice, MD, MSc; Matthew W. Semler, MD, MSc.
2019 2019	•	Multimorbidity Explorer A shiny app for exploring EHR and biobank data RStudio::conf 2019 • Contributed Poster. Authored with Yaomin Xu.
2019 2019	•	Taking a network view of EHR and Biobank data to find explainable multivariate patterns Vanderbilt Biostatistics Seminar Series • University wide seminar series.
2019		 Patient-specific risk factors independently influence survival in Myelodysplastic Syndromes in an unbiased review of EHR records Under-Review (copy available upon request.) Bayesian network analysis used to find novel subgroups of patients with Myelodysplastic Syndromes (MDS). Analysis done using method built for my dissertation.

2019	•	Patient specific comorbidities impact overall survival in myelofibrosis Under-Review (copy available upon request.)
		 Bayesian network analysis used to find robust novel subgroups of pa- tients with given genetic mutations. Analysis done using method built for my dissertation.
2018 	•	R timelineViz: Visualizing the distribution of study events in longitudinal studies
2018		Under-Review (copy available upon request.)
		 Authored with Alex Sunderman of the Vanderbilt Department of Epidemiology.
2017	•	Continuous Classification using Deep Neural Networks
2017		Vanderbilt Biostatistics Qualification Exam
2017		 Review of methods for classifying continuous data streams using neural networks
		 Successfully met qualifying examination standards
2015 I	•	Asymmetric Linkage Disequilibrium: Tools for Dissecting Multiallelic LD
2015		Journal of Human Immunology
		• Authored with Richard Single, Vanja Paunic, Mark Albrecht, and Martin Maiers.
2015	•	An Agent Based Model of Mysis Migration
2015		International Association of Great Lakes Research Conference
2015		\cdot Authored with Brian O'Malley, Sture Hansson, and Jason Stockwell.
2015	•	Declines of Mysis diluviana in the Great Lakes
 2015		Journal of Great Lakes Research
2013		

• Authored with Peter Euclide and Jason Stockwell.