JOSHUAWHITE

I'm an aspiring Cognitive & Data Scientist who last year finished his Honours in Psychology at the University of Melbourne. I have considerable experience as a research assistant at all levels of the scientific process. I have a particular interest in using computational methods and complex data analysis to better understand the mind, and have experience doing so with a variety of programs / programming languages. Previously, I worked as a litigation lawyer.



EDUCATION

2020 2015

PhD. Candidate. Biostatistics

Vanderbilt University

Nashville, TN

- · Focused on network models & interactive visualization platforms for electronic health records data
- · University Graduate Fellow

2015 2011

B.S., Mathematics, Statistics (minor C.S.)

University of Vermont

Burlington, VT

· Thesis: An agent based model of Diel Vertical Migration patterns of Mysis diluviana



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 clinically relevant patterns in data.

2018 2017 Data Science Researcher

Data Science Lab

Q Johns Hopkins University

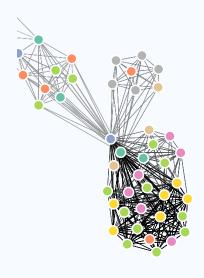
- · Building R Shiny applications in the contexts of wearables and statistics
- · Work primarily done in R Shiny and Javascript (node and d3js).

2015 2013

Undergraduate Researcher

Rubenstein Ecosystems Science Laboratory

- University of Vermont
- · 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.)



View this CV online with links at newgoldenhammer.com/cv/

CONTACT



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- github.com/WhiteJP
- **o** newgoldenhammer.com

https://www.linkedin.com/in/joshu p-white/

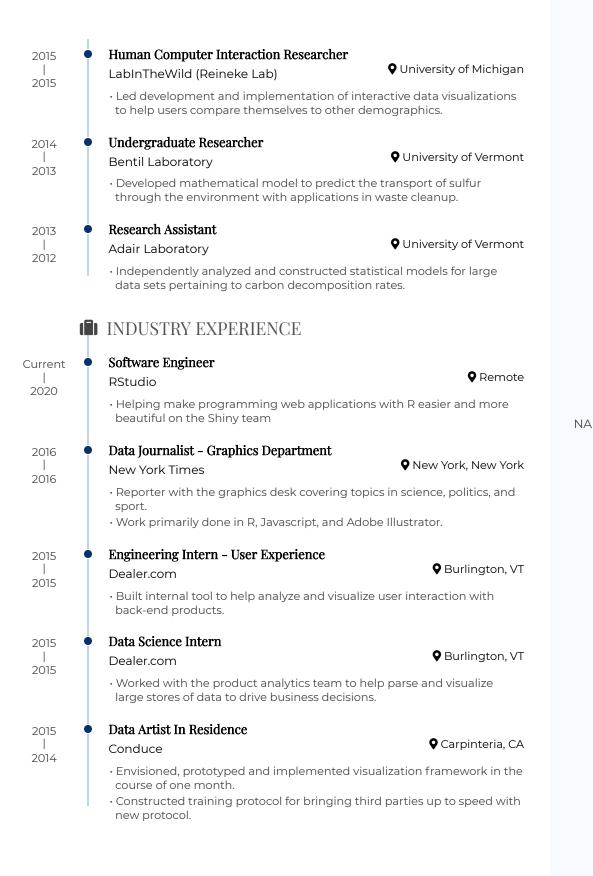
LANGUAGE SKILLS

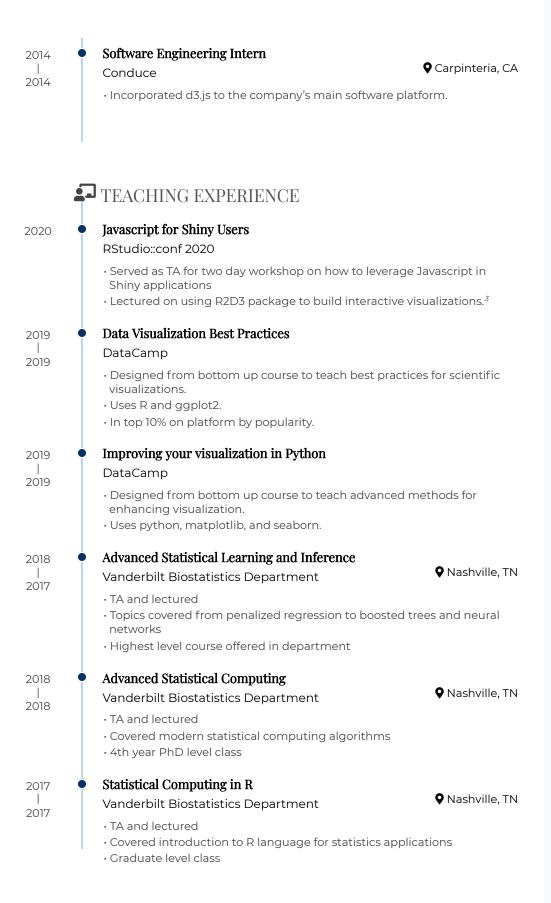
R	
Python	
MATLAB	
HTML	
Javascript	
CSS	

Made with the R package pagedown.

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

Last updated on 2021-01-28.





NA

✓ SELECTED DATA SCIENCE WRITING

2019 • Using AWK and R to Parse 25tb⁵

LiveFreeOrDichotomize.com

- · Story of parsing large amounts of genomics data.
- · Provided advice for dealing with data much larger than disk.
- · Reached top of HackerNews.

Classifying physical activity from smartphone data⁶

RStudio Tensorflow Blog

- Walk through of training a convolutional neural network to achieve state of the art recognition of activities from accelerometer data.
- · Contracted article.

2018 • The United States of Seasons⁷

LiveFreeOrDichotomize.com

- GIS analysis of weather data to find the most 'seasonal' locations in United States
- Used Bayesian regression methods for smoothing sparse geospatial data.

2017 • A year as told by fitbit⁸

LiveFreeOrDichotomize.com

- Analyzing a full years worth of second-level heart rate data from wearable device.
- · Demonstrated visualization-based inference for large data.

MCMC and the case of the spilled seeds

LiveFreeOrDichotomize.com

- Full Bayesian MCMC sampler running in your browser.
- · Coded from scratch in vanilla Javascript.

The Traveling Metallurgist¹⁰

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.

■ SELECTED PRESS (ABOUT)

2017 | 2017

2017

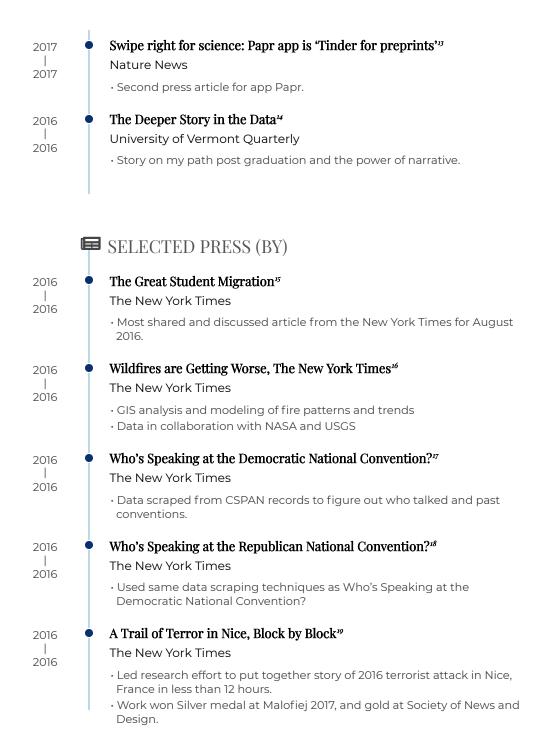
2018

Great paper? Swipe right on the new 'Tinder for preprints' app"

Science

 \cdot Story of the app Papr $^{\rm IZ}$ made with Jeff Leek and Lucy D'Agostino McGowan.

NA





■ 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 research

2020

Stochastic Block Modeling in R, Statistically rigorous clustering with rigorous code20

RStudio::conf 2020

- · Invited talk about new sbmR package²¹.
- · 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 PheWAS²²

Bioinformatics

- · Manuscript detailing application for the exploration of multimorbidity patterns in PheWAS analyses
- · See landing page²³ for more information.

2019 2019

Charge Reductions Associated with Shortening Time to Recovery in Septic Shock²⁴

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 data²⁵

RStudio::conf 2019

· Contributed Poster. Authored with Yaomin Xu.

2019 2019

Taking a network view of EHR and Biobank data to find explainable multivariate patterns²⁶

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.

Patient specific comorbidities impact overall survival in myelofibrosis 2019 Under-Review (copy available upon request.) · Bayesian network analysis used to find robust novel subgroups of patients with given genetic mutations. · Analysis done using method built for my dissertation. R timelineViz: Visualizing the distribution of study events in longitudinal 2018 studies 2018 Under-Review (copy available upon request.) · Authored with Alex Sunderman of the Vanderbilt Department of Epidemiology. Continuous Classification using Deep Neural Networks²⁷ 2017 Vanderbilt Biostatistics Qualification Exam 2017 · Review of methods for classifying continuous data streams using neural networks · Successfully met qualifying examination standards Asymmetric Linkage Disequilibrium: Tools for Dissecting Multiallelic LD 2015 Journal of Human Immunology 2015 · Authored with Richard Single, Vanja Paunic, Mark Albrecht, and Martin Maiers. An Agent Based Model of Mysis Migration²⁸ 2015 International Association of Great Lakes Research Conference 2015 · Authored with Brian O'Malley, Sture Hansson, and Jason Stockwell. 2015 Declines of Mysis diluviana in the Great Lakes Journal of Great Lakes Research



1: NA

2015

- 2: NA
- 3: http://nickstrayer.me/js4shiny_r2d3/slides
- 4: NA
- 5: https://livefreeordichotomize.com/2019/06/04/using_awk_and_r_to_parse_25tb/
- 6. https://blogs.rstudio.com/tensorflow/posts/2018-07-17-activity-detection/

· Authored with Peter Euclide and Jason Stockwell.

- 7: https://livefreeordichotomize.com/2018/02/12/the-united-states-of-seasons/
- 8: https://livefreeordichotomize.com/2017/12/27/a-year-as-told-by-fitbit/
- 9. https://livefreeordichotomize.com/2017/10/14/mcmc-and-the-case-of-the-spilled-seeds/
- 10: https://livefreeordichotomize.com/2017/09/25/the-traveling-metallurgist/
- 11: https://www.sciencemag.org/news/2017/06/great-paper-swipe-right-new-tinder-preprints-app
- 12: https://jhubiostatistics.shinyapps.io/papr/

- 13: https://www.nature.com/news/swipe-right-for-science-papr-app-is-tinder-for-preprints-1.22163
- 14: https://www.uvm.edu/uvmnews/news/deeper-story-data
- 15: https://www.nytimes.com/interactive/2016/08/26/us/college-student-migration.html?smid=pl-share
- 16. https://www.nytimes.com/interactive/2016/07/25/us/wildfire-seasons-losangeles.html
- 17: https://www.nytimes.com/2016/07/26/upshot/democrats-may-not-be-unified-but-their-convention-speakers-are.html
- 18: https://www.nytimes.com/2016/07/19/upshot/whos-not-speaking-how-this-republican-convention-differs.html?smid=pl-share
- 19: https://www.nytimes.com/interactive/2016/07/14/world/europe/trail-of-terror-france.html
- 20: http://nickstrayer.me/rstudioconf_sbm
- 21: https://tbilab.github.io/sbmR/
- 22. https://academic.oup.com/bioinformatics/advance-article-abstract/doi/10.1093/bioinformatics/btaa870/5922817?redirectedFrom=fulltext
- 23. https://prod.tbilab.org/phewas_me_info/
- 24. https://www.ncbi.nlm.nih.gov/pubmed/30419234
- 25. http://nickstrayer.me/rstudioconf19_me-poster/
- 26: http://nickstrayer.me/biostat_seminar/
- 27: http://nickstrayer.me/qualifying_exam/
- 28: https://www.semanticscholar.org/paper/An-Agent-Based-Model-of-the-Diel-Vertical-Migration-Strayer-Stockwell/40493c78e8ecf22bd882d17ec99fd913ec4b9820