Matthew N. Bernstein

http://mbernste.github.io

EDUCATION

University of Wisconsin – Madison

Madison, WI

Email: matthewb@cs.wisc.edu

Ph.D. in Computer Sciences

Expected 2019

o Advisor: Colin Dewey

• Thesis: Computational methods for transcriptome-based cellular phenotyping

University of Wisconsin – Madison

Madison, WI

M.S. in Computer Sciences

Dec. 2015

University of Notre Dame

South Bend, IN

B.S. in Computer Science; Magna Cum Laude

May 2013

EXPERIENCE

University of Wisconsin - Madison

Madison, WI

Research Assistant under Prof. Colin Dewey

Aug. 2014 - Present

- Researching and developing novel computational methods for extracting knowledge from large, public repositories of biological sequencing data
- Mentor undergraduate students working in the lab

Amazon Seattle, WA

Software Development Engineering Intern

Summer 2014

- Designed, implemented, and launched auto-complete search-suggestions for the Amazon Local website's search bar. Search suggestions are served as the user is typing a query
- Designed, implemented, and launched an offline system that uses previous customer search data to pre-compute an index of search suggestions

University of Wisconsin - Madison

Madison, WI

Teaching Assistant (Lecturer)

Aug. 2013 - May 2014

- Lecturer to ~30 students in CS 302 Introduction to Programming
- Created assignments, quizzes, and lesson plans for my class
- \circ Designed two programming projects that all \sim 700 students enrolled in CS 302 were required to complete

Amazon New York, NY

Software Development Engineering Intern

Summer 2013

- Improved Amazon Posts a tool that allows brands to create short social messages that appear on various feeds across Amazon websites
- Revamped the Amazon Posts management UI to include analytics that inform brands on how successfully each post has reached customers and promoted purchases
- o Improved the customer facing look and feel of the Amazon Posts feeds

Space and Naval Warfare Systems Command (SPAWAR)

San Diego, CA

Research Intern

 $Summer\ 2012$

- Researched applications of machine learning for the task of determining political and group affiliation of anonymous internet authors
- Built a prototype application for tagging public web content that may be of interest to intelligence analysts

AWARDS AND FELLOWSHIPS

- NIH/BD2K Young Investigator Travel Scholarship, International Conference on Intelligent Systems for Molecular Biology (ISMB), Chicago, IL, July 6-10, 2018
- Best Plenary Talk, National Library of Medicine Informatics Training Conference, San Diego, CA, June 6, 2017
- Awarded three year, NLM funded traineeship through the Computation and Informatics in Biology and Medicine training program between Feb. 2015 Feb. 2018
- University Housing Honored Instructor Award. University of Wisconsin-Madison, Fall 2013
- Tau Beta Pi Membership. University of Notre Dame, Fall 2011
- Upsilon Pi Epsilon Membership. University of Notre Dame, Fall 2011

PEER-REVIEWED PUBLICATIONS

• Bernstein, M.N., Doan, A., Dewey, C.N. (2017). MetaSRA: normalized human sample-specific metadata for the Sequence Read Archive. *Bioinformatics*, 33(18), 2914–2923.

INVITED TALKS

•	International Conference on Intelligent Systems for Molecular Biology (ISMB) MetaSRA: normalized human sample-specific metadata for the Sequence Read Archive	Chicago, IL July 7, 2018
•	Center for Predictive Computational Phenotyping Annual Retreat Cellular phenotyping with mass, heterogeneous transcriptomic data	Madison, WI May 31, 2018
•	National Library of Medicine Informatics Training Conference MetaSRA: normalized human sample-specific metadata for the Sequence Read Archive	San Diego, CA June 6, 2017
•	Center for Predictive Computational Phenotyping Annual Retreat MetaSRA: normalized human sample-specific metadata for the Sequence Read Archive	Madison, WI June 1, 2017

POSTER PRESENTATIONS

•	RNA-Seq Summit	San Francisco, CA
	MetaSRA: normalized human sample-specific metadata for the Sequence Read Archive	April 26-27, 2017
•	National Library of Medicine Informatics Training Conference	Columbus, OH
	Standardizing sample-specific metadata in the Sequence Read Archive	June 27-28, 2016

SERVICE

•	Integrated Biological Sciences Summer Research Program	Madison, WI
	Mentor	Summer 2015, 2016

• Co-mentored undergraduate students' summer research projects with Prof. Colin Dewey

Computer Sciences Graduate Student Welcome Weekend

Madison, WI Spring 2014, 2015, 2016

- Planned the department's prospective student visit weekend
- Chaired the committee in Spring 2015

Committee member, Committee chair

Scratch Computer Programming Club at Stephen's Point Elementary Club Leader Madison, WI Spring 2015

 \circ Led an after school computer science club for 4th and 5th grade students

TECHNICAL SKILLS

- Software Development: Python (strong), Java (strong), C/C++ (familiar), JavaScript (familiar), HTML, CSS, SQL, MongoDB, Git, Numpy, Matplotlib, Scikit Learn
- Computer science & Machine Learning: Probabilistic Modeling, Dimensionality Reduction, Named Entity Recognition, Knowledge Representation
- Bioinformatics Topics: RNA-seq, Transcriptome Quantification, Biomedical Text Mining