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

- Developing novel computational and statistical approaches for extracting knowledge from large, heterogeneous repositories of gene expression 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
- Designed two programming projects that all ~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

 $Summer\ 2012$

- Developed a machine learning based solution 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 Sci	ences 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

• 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