Shilpa Nadimpalli Kobren

□ +1 978-407-8042 • ☑ shilpa_kobren@hms.harvard.edu • ⑤ http://shilpakobren.com

Education

- 2018 **Ph.D. in Computer Science** Princeton University
 - Thesis: Detecting and Analyzing Variation in Protein Interaction Interfaces

Advisor: Mona Singh

- 2013 M.Sc. in Computer Science Princeton University
- 2011 **B.Sc. in Biology & Computer Science**, summa cum laude Tufts University

Thesis, Awarded Highest Honors: Correcting Protein Multiple Structural Alignments by Sequence Peeking

Advisor: Lenore Cowen

Awards & Honors

- 08/2015 **Siebel Scholar, Class of 2016** awarded annually for academic excellence and demonstrated leadership to 93 top students from the world's leading graduate schools
- 09/2011–09/2015 NSF Graduate Research Fellowship for graduate study in STEM disciplines
- 09/2011–09/2015 Princeton Gordon Wu Fellowship for engineering graduate study
 - 05/2011 **Tufts James Schmolze Award** for top undergraduate achievement in computer science
 - 05/2010 Google Anita Borg Memorial Scholarship for women in computer science
- 08/2007–05/2011 Tufts Neubauer Scholarship for aptitude in research; grants to fully cover all four years of

undergraduate student loans (1 in 10 of ~1500)

- 04/2011 Tufts Class of 1882 Prize Scholarship for intellectual leadership (1 in 4 of ~5000)
- 04/2010 Tufts Audrey Butvay Gruss Award for scientific achievement (1 in 4 of ~5000)
- 06/2009–10/2009 CRA-W/CDC Distributed Research Experiences for Undergraduates Fellowship
- 09/2008-05/2009 CRA-W Multidisciplinary Research Opportunities for Women Fellowship
- 06/2008–09/2008 NSF/NIH Southern California Bioinformatics Summer Institute Fellowship
 - 05/2006 Massachusetts State Science Fair, 1st Place Award

Publications

- 2019 **SN Kobren**, B Chazelle, M Singh. (2019). "An integrative approach to identify preferentially altered interactions in human cancers." *Manuscript in submission*.
 - **SN Kobren**, M Singh. (2019). "Systematic domain-based aggregation of protein structures highlights DNA-, RNA- and other ligand-binding positions." *Nucleic Acids Research*. 47: 582–593.
- **S Nadimpalli**, AV Persikov, M Singh. (2015). "Pervasive variation of transcription factor orthologs contributes to regulatory network divergence." *PLoS Genetics*. 11: e1005011.
- N Daniels, **S Nadimpalli**, L Cowen. (2012). "Formatt: Correcting protein structural alignments by incorporating sequence alignment." *BMC Bioinformatics*. 13: 259–266.
- **S Nadimpalli***, N Daniels*, L Cowen. (2011). "Formatt: Correcting protein structural alignments by sequence peeking." *Proceedings of the 2011 ACM Conference on Bioinformatics, Computational Biology, and Biomedicine*. 2: 315–319.
- J Rieffel, F Sauders, **S Nadimpalli**, H Zhou, S Hassoun, J Rife, B Trimmer. (2009). "Evolving soft robotic locomotion in PhysX." *Proceedings of the 2009 ACM Conference on Genetic and Evolutionary Computation*. 11: 2499–2504.

Research & Work Experience

08/2018 – Present	Postdoctoral Research Fellow , Harvard Medical School • <i>Boston, MA</i> Developing translational bioinformatics applications that combine insights from genomics data and electronic health records to improve diagnoses of rare diseases under Dr. Isaac S. Kohane
03/2012–06/2018	Graduate Research Assistant, Princeton University • Princeton, NJ "Detecting and Analyzing Variation in Protein Interaction Interfaces" Investigated the evolution, natural variation, and disease-related mutation of cellular networks through analysis of protein interaction interfaces under Prof. Mona Singh
	Undergraduate Research Assistant, Tufts University • Medford, MA
01/2010-05/2011	"Formatt: Correcting Protein Multiple Structural Alignments by Sequence Peeking" Improved the Matt protein structural aligner by incorporating a sequence alignment correction step and validating using objective measures under Prof. Lenore J. Cowen
09/2008-05/2009	"Modeling the Neuro-Mechanical Control of a Soft-Tissue Organism" Evolved a solution representing the firing of neurons for muscle contraction using genetic algorithms to successfully elicit motion in a SoftBot under Prof. Soha Hassoun
06/2009–10/2009	Summer Intern, Bioinformatics Dept, Virginia Tech • Blacksburg, VA "Cataloging Animal Retrocopies for Retrotransposon Model" Developed a genome annotation independent algorithm to identify and classify retrocopies and chimeric genes given a transcriptome under Prof. Liqing Zhang
06/2008-09/2008	Summer Intern, Biochemistry Dept, University of California, Los Angeles • Westwood, CA "Discovery of Novel Metabolic Types of Bacterial Microcompartments" Implemented a pipeline to analyze oceanic metagenomic sequencing data for bacterial microcompartment (BMC) shell proteins to identify potentially novel metabolic BMC types under Prof. Todd Yeates
05/2007–08/2007	Summer Intern, Preclinical Dept , Shire Pharmaceuticals • <i>Cambridge & Lexington, MA</i> Set up a secure database containing clinical and research reports for drugs in production or under testing.
06/2006-09/2006	Summer Intern, Microbiology Dept, Univ. of Massachusetts Medical School • Worcester, MA "Identification of Genes Involved in Expansion of Chromosomal Repeat in E. Coli" Developed a series of knockout bacterial strains and tested viability of bacteria and amplification of a specific chromosomal repeat under Prof. Anthony Poteete

Teaching & Advising Experience

leaching & Advising Experience		
11/15/2018	Guest Lecturer in Computational Biology , Tufts University • <i>Medford, MA</i> Gave 1-hour lecture on DNA motif finding algorithms to undergraduate and graduate students in <i>CS167</i> .	
01/2016-09/2017	Coadviser for First-Year Graduate Students, Princeton University • Princeton, NJ Fall 2016, Spring 2016 & Spring 2017 Coadvised three Ph.D. students in Computational Biology and Computer Science on separate projects related to protein domains, protein binding and cancer	
06/2013-07/2013	Coadviser for Undergraduate Students, Princeton University • Princeton, NJ Summer 2013 Coadvised three undergraduates on a summer project through the Summer Programming Experiences program to build a secure voting mobile application	
09/2012–05/2013	Assistant Instructor, Princeton University • Princeton, NJ Fall 2012 & Spring 2013 Taught precepts twice weekly for COS126: General Computer Science, held office hours (~10 hours/week), graded assignments and proctored midterm and final exams. *Top rated (4.5/5 on average) by students for all precepts taught over both semesters.	
09/2009-05/2010	Teaching Assistant , Tufts University ● <i>Medford, MA</i> Fall 2009 & Spring 2010 Attended lectures, assisted in labs, held office hours (~10-18 hours/week) and graded assignments for two courses CS40: Machine Structure and Assembly Language Programming and CS80: Programming Languages.	
02/2009-06/2009	High School Bioinformatics Instructor , Somerville High School • <i>Somerville</i> , <i>MA</i> Analyzed soil metagenomic data to discover new microbes and led four spring seminars for students about gene sequencing and alignment algorithms. Project funded via an HHMI ARRAYS grant.	

Presentations

Invited Talks

10/25/2018	'Rising Stars in Computer Science' Symposium at Tufts University, Medford, MA
03/08/2018	Open Insights Seminar Series at Harvard Medical School, Dept. of Biomedical Informatics, Boston, MA
01/09/2018	Seminar at the Flatiron Institute, Center for Computational Biology (CCB), New York City, NY
11/03/2017	Lewis-Sigler Institute for Integrative Genomics 2017 Retreat, Princeton, NJ
09/26/2017	Data Science Seminar Series at the University of Massachusetts, Amherst, MA
09/13/2017	'Rising Stars in Data Science' Symposium at the University of Chicago, Chicago, IL
01/06/2017	Princeton Area Alumni Association First Friday Series, Princeton, NJ
01/06/2016	Icahn Institute Think & Drink Symposium, Princeton, NJ
	Platform Presentations
08/02/2017	NY Area Meeting in Quantitative Biology, Cold Spring Harbor, NY
04/02/2011	Tufts 13th Annual Undergraduate Research Symposium, Medford, MA
	Posters
11/15/2015	RECOMB/ISCB Conference on Regulatory and Systems Genomics, Philadelphia, PA
03/20/2014	Systems Biology Meeting: Global Regulation of Gene Expression, Cold Spring Harbor, NY
04/09/2011	3rd Annual New England Undergraduate Computing Symposium, Cambridge, MA
10/09/2009	Midwest Women in Computing Conference, Chicago, IL
	Panels
07/23/2019	"Doctoral Career Trajectories" at Harvard DBMI Summer Institute in Biomedical Informatics, Boston, MA
08/22/2017	"Women in Bioinformatics" at ACM Conference on Bioinformatics, Computational Biology, and Biomedicine, Boston, MA
10/02/2015	"Pursuing Graduate Studies in Computer Science" at Princeton University Computer Science Dept., Princeton, NJ
03/28/2013	"Maximizing the Utility of Office Hours" at Princeton University McGraw Teaching Center, Princeton, NJ
04/09/2011	"Graduate School Application Process and Visit Experience" at 3rd Annual New England Undergraduate Computing Symposium, <i>Boston, MA</i>

Service

Conferences

01/2019-Present	Pacific Symposium on Biocomputing 2019, Session Co-Chair and referee, Big Island of Hawaii
01/2017-08/2017	ACM Bioinformatics & Computational Biology 2017 Student Activity Co-Chair and referee, Boston, MA
05/2017, 05/2019	RECOMB 2017 & 2019 referee, Hong Kong & Washington, DC
07/2010, 07/2014	ISMB 2010 & 2014 Student Volunteer and referee, Boston, MA
	Leadership
10/2018-Present	Undiagnosed Diseases Network, Tool Building Coalition Working Group, Primary Organizer
05/2012-02/2015	Princeton Graduate Student Government, Computer Science Representative, Events Board Member
07/2012-05/2014	Princeton Jewish Graduate Students and Young Professionals, President
	Mentoring and Outreach
01/2019-Present	Harvard Women in STEM (WiSTEM) Mentoring Program, Undergraduate Student Mentor
09/2011-05/2018	Princeton Graduate Women in Science and Engineering, Mentoring Program Participant
08/2011-Present	Tufts Alumni Admissions Program, Applicant Interviewer
09/2008-05/2011	Tufts Admissions Office, April Open House Host, Voices Host, Speaker at Engineering Open House
09/2007-05/2011	Tufts Association for Computing Machinery, Women, Mentor & Outreach Contact

Last updated: 13-Aug-2019