Shilpa Nadimpalli Kobren

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Currently seeking a position as a postdoctoral researcher in the New York City or Boston areas.

Education

Princeton University, School of Engineering • Princeton, NJ

Ph.D. in Computer Science, Advisor: Prof. Mona Singh
Thesis: "Detecting and Analyzing Variation in Protein Interactions"

June 2013 M.A. in Computer Science

May 2011 Tufts University, School of Arts & Sciences • Medford, MA
B.S. in Biology & Computer Science, summa cum laude
Senior Thesis, Awarded Highest Honors: "Formatt: Correcting Protein Multiple
Structural Alignments by Sequence Peeking"

Awards, Honors, & Fellowships

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's Graduate Research Fellowship for graduate study in STEM disciplines
09/2011 - 09/2015	Princeton's Gordon Wu Fellowship for engineering graduate study
08/2010 - 05/2011	Google's Anita Borg Memorial Scholarship for women in computer science
08/2007 - 05/2011	Tufts' Neubauer Scholarship for aptitude in research (1 in 10 of ~1350)
05/2011	Tufts' James Schmolze Award for excellence in computer science (1 of 42)
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's Distributed Research Experiences for Undergrads Fellowship
09/2008 - 05/2009	CRA-W's Multidisciplinary Research Opportunities for Women Fellowship
06/2008 - 09/2008	NSF/NIH's Southern California Bioinformatics Summer Institute Fellowship

Peer-Reviewed Publications

- **S Nadimpalli**, AV Periskov, 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 Saunders, **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 (GECCO '09)*. 11: 2499-2504.

Last updated: 17-Apr-2017

Research & Work Experience

03/2012 – Present	Princeton University, Dept. of Computer Science • Princeton, NJ
	Investigating the evolution, natural variation, and cancer-related mutation of gene regulatory
	networks through analysis of protein interaction interfaces under Prof. Mona Singh

01/2010 – 05/2011 **Tufts University,** Dept. of Computer Science • Medford, MA "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 Cowen.

Virginia Tech, Dept. of Computer Science • Blacksburg, VA

"Cataloging Animal Retrocopies for Retrotransposition Model"

Developed a genome annotation independent algorithm to identify and classify retrocopies and chimeric genes in animal genomes given a transcriptome as input under Prof. Liqing Zhang.

09/2008 – 05/2009 **Tufts University,** Depts. of Comp. Sci., Biology, & Biomedical Eng. • Medford, MA "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.

06/2008 – 09/2008 UCLA, Dept. of Biochemistry • Westwood, CA

"Discovery of Novel Metabolic Types of Bacterial Microcompartments"

Implemented a pipeline for metagenomes to analyze potential proteins within bacterial microcompartments (BMCs) to identify novel metabolic BMC types under Prof. Todd Yeates.

05/2007 – 08/2007 Shire Pharmaceuticals, Preclinical Dept. • Cambridge & Lexington, MA Set up a secure database containing all clinical and research reports for all drugs currently produced or under testing.

06/2006 – 09/2006 **UMass Medical School,** Dept. of Molecular Genetics • 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.

Posters, Presentations, & Panels

Aug. 20-23, 2017	ACM Conference on Bioinformatics, Comp. Bio. & Biomedicine (BCB) • Boston, MA
Jan. 6, 2017	Student Activity Co-Chair Princeton Area Alumni Association First Friday Series • Princeton, NJ
	<i>Talk:</i> "Structure-informed approach to discovering perturbed interaction interfaces in cancer"
Jan. 6, 2016	Icahn Institute Think & Drink Symposium • Princeton, NJ
	Talk: "Pervasive variation of TF orthologs contributes to regulatory network divergence"
Mar. 18-22, 2014	
	Poster: "Pervasive binding specificity variation Cys ₂ -His ₂ zinc finger orthologs suggests trans
	mutations as major drivers of regulatory network divergence"
Apr. 9, 2011	3rd Annual New England Undergraduate Computing Symposium • Medford, MA
	Panelist: Graduate School Application Process and Visit Experience
	Poster: "Formatt: Correcting Protein Structural Alignments by Sequence Peeking"
Apr. 2, 2011	Tufts 13 th Annual Undergraduate Research Symposium • Medford, MA
	Talk: "Formatt: Correcting Protein Structural Alignments"
July 8-13, 2010	Internat'l Conference on Intelligent Systems for Molecular Bio. (ISMB) • Boston, MA
	Student Volunteer
Apr. 17, 2010	2nd Annual New England Undergraduate Computing Symposium • Boston, MA
	Poster: "Improving Matt Structural Alignments by Considering Sequence Homology"
Oct. 9-10, 2009	
	Poster: "Cataloging Animal Retrocopies: An Annotation-Independent Methodology"

Teaching Experience

01/2016 – 02/2017	Princeton University, Dept. of Computer Science • Princeton, NJ Mentor for first-year graduate students (Spring 2016 & Fall 2016) Mentored incoming Quantitative and Computational Biology graduate student — "Identifying functional protein domain positions using population variation data" Mentored rotating Computer Science graduate student — "Framework for structural integration of non-synonymous mutations in protein binding pockets"
06/2013 – 07/2013	Mentor for Summer Programming Experiences (Summer 2013) Co-advised three undergraduates on a summer project to build a secure voting mobile app
09/2012 – 05/2013	Assistant Instructor for COS126: General Computer Science (Fall 2012 & Spring 2013) Teach precepts twice weekly, hold office hours, and grade assignments and exams.
09/2009 – 05/2010	Tufts University, Dept. of Computer Science • Medford, MA <i>TA for CS40: Machine Structure and Assembly Language Programming (Fall 2009) TA for CS80: Programming Languages (Spring 2010)</i> Attended lectures, assisted in labs, held office hours (~10-18 hrs/wk), and graded assignments.
02/2009 – 06/2009	Tufts University, Dept. of Chemistry • Medford, MA HHMI ARRAYS Project Programmer & Instructor Analyzed Illumina sequencing output to discover new soil microbes, and taught bioinformatics seminars to Somerville High students about gene sequencing, alignment algorithms, and BLAST.

Extracurricular Activities & Outreach

09/2011 – Present	Princeton Grad. Women in Science and Engineering, Mentoring Program Participant
08/2011 - Present	Tufts Alumni Admissions Program (TAAP), Applicant Interviewer
05/2012 - 02/2015	Princeton Grad. Student Government, CS Representative, Events Board member
07/2012 - 05/2014	Princeton Jewish Grad. Students & Young Professionals, President, Coordinator
10/2009 - 05/2011	Tufts Computer Science Reading Group, Member & Coordinator
09/2008 - 05/2011	Tufts Admissions Office, April Open House / Voices Host, Speaker at Engineering OH
09/2007 - 05/2011	Tufts Association for Computing Machinery, Women, Mentor & Outreach Contact

Last updated: 17-Apr-2017