# Shilpa Nadimpalli Kobren

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

#### **Education**

- exp. 2017 **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**

- **SN Kobren**, M Singh. (2017). "Structure-informed approach to discovering perturbed interaction interfaces in cancer." *Manuscript in preparation*.
  - **SN Kobren**, M Singh. (2017). "Structure-based assessment of domain binding potential." *Manuscript in preparation*.
- **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*. 2: 315–319.

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## **Research & Work Experience**

03/2012–Present Princeton University, Graduate Research Assistant • Princeton, NJ

"Detecting and Analyzing Variation in Protein Interaction Interfaces"

Investigating the evolution, natural variation, and disease-related mutation of gene regulatory networks through analysis of protein interaction interfaces under Prof. Mona Singh

Tufts University, Undergraduate Research Assistant • 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 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 Virginia Tech, Summer Intern • 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 University of California, Los Angeles, Biochemistry Department Summer Intern • 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 Department Summer Intern • Cambridge & Lexington, MA

Set up a secure database containing clinical and research reports for drugs in production or under testing.

06/2006–09/2006 University of Massachusetts Medical School, Summer Intern • 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 & Mentoring Experience**

01/2016-02/2017 Graduate Student Mentor, Princeton University • Princeton, NJ

Spring 2016 Quantitative & Computational Biology graduate student project: "Identifying functional

protein domain positions using population variation data"

Fall 2016 Computer Science graduate student project: "Framework for structural integration of

nonsynonymous mutations in protein binding pockets"

06/2013-07/2013 Undergraduate Student Mentor, Princeton University • Princeton, NJ

Summer 2013 Co-advised 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 ( $\sim$ 10 hours/week), graded assignments and proctored midterm and final exams.

\*Top rated (4.95/5 on average) by students for all precepts taught over both semesters.

09/2009–05/2010 **Teaching Assistant**, Tufts University • Medford, MA

Fall 2009 & Spring 2010 Attended lectures, assisted in labs, held office hours ( $\sim$ 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 • Somerville, MA

Analyzed *Illumina* output to discover new soil microbes and led seminars for Somerville High students about gene sequencing, alignment algorithms, and BLAST. Project funded via a Tufts University Chemistry

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Department HHMI ARRAYS grant.

# **Presentations**

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01/06/2017	Structure-informed approach to discovering perturbed interaction interfaces in cancer, <i>Princeton Area Alumni Association First Friday Series, Princeton, NJ</i>
01/06/2016	Pervasive variation of TF orthologs contributes to regulatory network divergence, Icahn Institute Think & Drink Symposium, Princeton, NJ
04/02/2011	Formatt: Correcting Protein Structural Alignments, Tufts 13th Annual Undergraduate Research Symposium, Medford, MA
	Posters
03/20/2014	Pervasive binding specificity variation of Cys <sub>2</sub> -His <sub>2</sub> zinc finger orthologs suggests trans mutations as major drivers of regulatory network divergence, <i>Systems Biology Meeting: Global Regulation of Gene Expression, Cold Spring Harbor, NY</i>
04/09/2011	Formatt: Correcting protein structural alignments by sequence peeking, 3rd Annual New England Undergraduate Computing Symposium, Boston, MA
04/17/2010	Improving Matt Structural Alignments by Considering Sequence Homology, 2rd Annual New England Undergraduate Computing Symposium, Boston, MA
10/09/2009	Cataloging Animal Retrocopies: An Annotation-Independent Methodology, Midwest Women in Computing Conference, Chicago, IL
	Panels
10/02/2015	Questions about Going to Graduate School in Computer Science, Department of Computer Science at Princeton University, Princeton, NJ
03/28/2013	Getting More Out of Office Hours, Princeton University McGraw Teaching Center, Princeton, NJ
04/09/2011	Graduate School Application Process and Visit Experience, 3rd Annual New England Undergraduate Computing Symposium, Boston, MA

## **Service & Outreach**

08/2017	<b>ACM BCB'17 Student Activity Co-Chair</b> , ACM Conference on Bioinformatics, Computational Biology, and Biomedicine, Boston, Massachusetts
05/2017	<b>RECOMB 2017 Sub-reviewer for submitted papers</b> , 21st Annual International Conference on Research in Computational Molecular Biology, Hong Kong
07/2010, 07/2015	<b>ISMB 2010 &amp; 2015 Student Volunteer</b> , International Conference on Intelligent Systems for Molecular Biology, Boston, Massachusetts
05/2012-02/2015	Princeton Graduate Student Government, Computer Science Representative, Events Board Member
09/2011-Present	Princeton Graduate Women in Science and Engineering, Mentoring Program Participant
08/2011-Present	Tufts Alumni Admissions Program, Applicant Interviewer
07/2012-05/2014	Princeton Jewish Graduate Students and 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 Host, Voices Host, Speaker at Engineering Open House
09/2007-05/2011	Tufts Association for Computing Machinery, Women, Mentor & Outreach Contact

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