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 \sim 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 Graduate Research Assistant, Princeton University • 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

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 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 • Cambridge & Lexington, MA

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 & Mentoring Experience

01/2016-02/2017 Mentor for First-Year Graduate Students, 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 Mentor for Undergraduate Students, 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 (4.5/5 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 (~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*-sequenced soil metagenomic data to discover new soil microbes, and led four spring seminars for Somerville High School students about gene sequencing, alignment algorithms, and BLAST.

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Project funded via a Tufts University Chemistry Department HHMI ARRAYS grant.

Presentations

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	Turko
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, <i>Icahn Institute Think & Drink Symposium, Princeton, NJ</i>
04/02/2011	Formatt: Correcting Protein Structural Alignments, Tufts 13th Annual Undergraduate Research Symposium, Medford, MA
	Posters
11/15/2015	Pervasive variation of transcription factor orthologs contributes to regulatory network divergence, RECOMB/ISCB Conference on Regulatory and Systems Genomics, Philadelphia, PA
03/20/2014	Pervasive binding specificity variation of Cys ₂ -His ₂ 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

Service & Outreach

08/2017	ACM BCB'17 Student Activity Co-Chair , ACM Conference on Bioinformatics, Computational Biology, and Biomedicine, Boston, Massachusetts
05/2017	RECOMB 2017 Sub-reviewer for submitted papers , 21st Annual International Conference on Research in Computational Molecular Biology, Hong Kong
07/2010, 07/2015	ISMB 2010 & 2015 Student Volunteer , 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

04/09/2011 Graduate School Application Process and Visit Experience, 3rd Annual New England Undergraduate

Computing Symposium, Boston, MA

References

Mona Singh Ph.D. Advisor	Lenore Cowen Undergraduate Research Advisor	Olga Troyanskaya Dissertation Committee Member
mona@cs.princeton.edu + 1 609-258-7059	cowen@cs.tufts.edu +1 617-627-5134	ogt@genomics.princeton.edu + 1 609-258-1749
Dept. of Computer Science and Lewis-Sigler Institute for Comparative Genomics Princeton University, Princeton, NJ	Dept. of Computer Science and Dept. of Mathematics Tufts University, Medford, MA	Dept. of Computer Science and Lewis-Sigler Institute for Comparative Genomics Princeton University, Princeton, NJ

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