# Hirak Sarkar

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#### Research Interest

I am interested in applying machine learning techniques such as statistical inference and deep learning to analyse and extract information from big data in the field of genomics, social and computer networks.

#### Education

## Ph.D in Computer Science

2014-2019

Statistical Inference in Biological Data, advisor: Prof. Rob Patro

Stony Brook University, NY

GPA: 3.99/4.00

## Masters of Technology (Computer Science)

2011-2013

Indian Statistical Institute  $1^{st}$  Class (Hons.)

# Bachelor of Technology (Computer Science and Engineering)

2007-2011

West Bengal University of Technology

GPA: 8.88/10

#### **Publications**

- 1. Towards selective-alignment: Bridging the accuracy gap between alignment-based and alignment-free transcript quantification, by <u>Hirak Sarkar\*</u>, Mohsen Zakeri\*, Laraib Malik, Rob Patro. [Submitted to Bioinformatics, 2017]
- 2. Quark enables semi-reference-based compression of RNA-seq data by <u>Hirak Sarkar</u> and Rob Patro [accepted Bioinformatics'17, impact factor: 7.307].
- 3. Fast, Lightweight Clustering of de novo Transcriptomes using Fragment Equivalence Classes by A Srivastava\*, <u>Hirak Sarkar\*</u>, Laraib Malik and Rob Patro (\* Joint first authors) [RECOMB-seq'16].
- 4. RapMap: A Rapid, Sensitive and Accurate Tool for Mapping RNA-seq Reads to Transcriptomes by A Srivastava, <u>Hirak Sarkar</u>, Nitish Gupta and Rob Patro [ISMB'16, acceptance rate: 17%].
- 5. Voronoi Game on Graphs (Extended version) by S. Bandyapadhyay, A. Banik, S. Das and H. Sarkar (in alphabetical order of surnames) Journal of Theoretical Computer Science [TCS'15].
- Voronoi Game on Graphs by S. Bandyapadhyay, A. Banik, S. Das and <u>H. Sarkar</u> (in alphabetical order of surnames) Seventh International Workshop on Algorithms and Computation. WALCOM'13.
- 7. Pufferfish: A fast graph-based indexing and query strategy for large genomic sequences by Fatemeh Almodaresi\*, <u>Hirak Sarkar\*</u>, Yi-Fei and Rob Patro, Poster presented in [WABI'17].
- 8. Joint probabilistic model for multiple steps of gene regulation by <u>Hirak Sarkar</u>, Yi-Fei Huang and Adam Siepel, Poster presented in **BioData'16**

## Professional Experience

- Simons Center for Quantitative Biology, Cold Spring Harbor Lab: Worked under the supervision of Prof. Adam Seipel from May, 2016 to July, 2016. We designed probabilistic graphical model to infer transcription and degradation rates from different assays such as GRO-seq and RNA-seq.
- Summer Assistantship '15,'17 with Prof. Rob Patro from May, 2015 to July, 2015. We worked on various problems ranging from

- Teaching Assistant for CSE549 (Computational Biology), CSE219 (Game Programming)
- Visiting Researcher at Advanced Computing & Microelectronics Unit, Indian Statistical Institute from October, 2013 to December, 2013. I worked on Computational Geometry and Graph Theory
- Junior Research Fellow in Department of Computer Science & Engineering at Indian Institute of Technology, Kharagpur (IIT) from July, 2013 to Sept, 2013. I was a member of Complex Network Engineering Group. I did TA-ship for Introductory Programming Course in that brief stint.

### Relevant Course Projects

• *IPID Header Survey:* We used IPID headers to estimate the load over different servers, sampled from alexa list. The main challenge of the project is to detect the wrapping pattern and navigate through the global vs local IPID counter. We also looked at the temporal pattern of network traffic for the different regional websites which shows interesting correlation with possible working load at the server end.

Instructor: Prof. Phillipa Gill

- Classification & Clustering of satellite images: We benchmarked different classification algorithms to cluster different kind of landmasses from infrared images taken by satellite.

  Instructor: Prof. C A Murthy
- Some Geometric and Combinatorial Properties of Binary Matrices Related to Discrete Tomography: Here we are trying to decompose an image matrix into matrices each having orthogonal convex polygon also known as Ferrer?s digraph. An operation could regenerate the original image from these matrices. The methods can be applied to image and data compression. (Masters dissertation) Advisor: Prof. Bhargab B Bhattacharya & Prof. Sandip Das
- GameSAT- A Structured Approach to Combine SLS SAT Solvers: Here we used several existing heuristic algorithms to mix up with each other in a customized probability to solve combinatorial hard problems encoded as SAT problems. We used UBCSAT framework for experimentation.

(B. Tech dissertation) Advisor: Ashiqur KhudaBukhsh, CMU

#### Awards and Honors

- Awarded Research Assistantship, SBU (2016-present)
- Awarded Special CS Chair Fellowship (of \$10000), SBU (2014-2015)
- Awarded NUS Research Scholarship, NUS (Jan'14-June'14)
- Awarded Post-graduate Scholarship by, Govt. of India. (2011-2013)
- Received First Prize for Software Competition (IEM), Calcutta.

#### **Programming Skills**

Python, C++,C

### Open Source Tools Used

Dendropy, BioNet (Comp Bio) NLTK, Scrapy, Scikit-learn, Stanford Parser, Pandas (Data Science)

# Relevant Coursework

- ullet Artificial Intelligence, Computational Biology, Analysis of Algorithms, Fundamental of Networks. (at SBU)
- Machine Learning & Pattern Recognition, Image Processing, Stochastic Process, Optimization Algorithms, Computer Graphics. (at *Indian Statistical Institute*)