Narmada Sambaturu

259, 9th cross, Tata Nagar, Kodigehalli, Bangalore - 560092, Karnataka, India

Website: https://narmadasambaturu.github.io/

Email: narmada.sambaturu@gmail.com

Research Interests Computational Biology, Systems Biology, Epidemiology, Immunotherapy, Multi-scale Modelling, Immune Response, Cancer Genomics, Host-pathogen Interactions.

Education

PhD, Interdisciplinary Mathematical Sciences, 2015 - 2021 (expected)

Mathematical and Computational Biology Stream,

IISc Mathematics Initiative,

Indian Institute of Science, Bangalore, India.

Supervisors: Prof. Nagasuma Chandra, Prof. N. Srinivasan.

Thesis (submitted): Multi-scale Modelling of Immune Response and Disease Spread: Methods and Applications.

Master of Science, School of Computing,

2012 - 2015

National University of Singapore, Singapore.

Supervisor: Prof. Wing-Kin Sung.

Thesis: Towards Handling Repeats in Genome Assembly.

Bachelor of Engineering, Computer Science and Engineering, 2005 - 2009

M.S.Ramaiah Institute of Technology, Bangalore, India.

Visvesvaraya Technological University.

Research Experience

• PhD Fellow,

2015 - 2021 (expected)

Interdisciplinary Mathematical Sciences, IISc Mathematics Initiative,

Indian Institute of Science, Bangalore, India.

Supervisors: Prof. Nagasuma Chandra, Prof. N. Srinivasan.

• Visitor, Winter 2016

Department of Applied Mathematics,

School of Mathematics,

University of Leeds, Leeds, UK

Research advisors: Prof. Carmen Molina-París, Prof. Grant Lythe.

• Junior Research Fellow,

Summer 2015

Indian Institute of Science, Bangalore, India. Research advisor: Prof. Nagasuma Chandra.

• Intern, 2006

Bioinformatics Centre,

Indian Institute of Science, Bangalore, India.

Research advisor: Prof. K. Sekar.

Work Experience Developer, Tata Consultancy Services

2009 - 2011

Technology Excellence Group,

Bangalore, India.

Teaching Experience

• Teaching Assistant for course Current trends in drug discovery. Indian Institute of Science, Bangalore, Karnataka, India.

Summer 2018

• Learning Enabler, Tata Consultancy Services. Common Initial Learning Program. Bidadi, Karnataka, India. 2010

Awards

- Best Student Paper Award, 2015 IEEE International Conference on Bioinformatics and Biomedicine (BIBM).
- One of 50 students selected from all over India for participation in a summer camp in Biotechnology at M.S.Swaminathan Research Foundation, Chennai, India (2003).
- All India Rank 25 in Secondary School Certificate (Std X) (2003).

Publications

- Narmada Sambaturu, Vaidehi Pusadkar, Sridhar Hannenhalli, Nagasuma Chandra, "PathExt: a general framework for path-based mining of omics-integrated biological networks." *Bioinformatics* 2020, doi:10.1093/bioinformatics/btaa941
- Narmada Sambaturu, Sumanta Mukherjee, Martín López-García, Carmen Molina-París, Gautam I. Menon, and Nagasuma Chandra. "Role of genetic heterogeneity in determining the epidemiological severity of H1N1 influenza." *PLoS Computational Biology* 14, no. 3 (2018): e1006069.
- 3. Narmada Sambaturu, Madhulika Mishra, and Nagasuma Chandra. "EpiTracer-an algorithm for identifying epicenters in condition-specific biological networks." *BMC genomics* 17, no. 4 (2016): 543.
- 4. Narmada Sambaturu, Madhulika Mishra, and Nagasuma Chandra. "EpiTracer-an algorithm for identifying epicenters in condition-specific biological networks." *Proceedings of the 2015 IEEE International Conference on Bioinformatics and Biomedicine (BIBM)*. IEEE Computer Society, 2015. (Best Student Paper)
- Narmada Sambaturu. "Towards handling repeats in genome assembly." MSc dissertation., 2014.
- Sridhar, Jayavel, Narmada Sambaturu, Radhakrishnan Sabarinathan, Hong-Yu Ou, Zixin Deng, Kanagaraj Sekar, Ziauddin Ahamed Rafi, and Kumar Rajakumar. "sR-NAscanner: a computational tool for intergenic small RNA detection in bacterial genomes." PLOS ONE 5, no. 8 (2010): e11970.
- 7. Annapurna P. Patil, Narmada Sambaturu, and Krittaya Chunhaviriyakul. "Convergence time evaluation of algorithms in MANETs." *International Journal of Computer Science and Information Security*, Vol. 5, No. 1, pp. 144-149, September 2009.

Workshop and Conference Presentations

- Narmada Sambaturu, Sumanta Mukherjee, Martín López-García, Carmen Molina-París, Gautam I Menon, and Nagasuma Chandra. (Talk) *Incorporating genetic heterogeneity into epidemic models for H1N1 influenza*. Mathematical and Statistical Explorations in Disease Modelling and Public Health, International Centre for Theoretical Sciences (ICTS), Bangalore, India, Jul 1 11 2019.
- Narmada Sambaturu, Madhulika Mishra, Rahul Metri and Nagasuma Chandra. An Algorithm for Identifying Druggable Targets Among Influential Mutations in Individual Cancer Patients. (Poster) Indo-US conference on Sculpting the future of medicine Gateway to the post-proteogenome era, at Advanced Centre For Treatment, Research And Education In Cancer (ACTREC), Mumbai, India, Dec 10 11 2018.
- Narmada Sambaturu and Nagasuma Chandra. OptiNeo an algorithm to optimise the number of neo-antigenic peptides for cancer immunotherapy. (Poster) Nature Big Data and Cancer Precision Medicine, Boston, Massachusetts, Oct 1 2 2018.
- Narmada Sambaturu, Sridhar Hannenhalli, and Nagasuma Chandra. (Poster) Cutting through the complexity of genomic data: A general method to identify candidate genes. RECOMB/ISCB Conference on Regulatory and Systems Genomics with DREAM Challenges, New York, NY, Nov 19 21 2017.

• Narmada Sambaturu, Sumanta Mukherjee, Martín López-García, Carmen Molina-París, Gautam I. Menon, and Nagasuma Chandra. (Talk) Role of genetic heterogeneity in determining the epidemiological severity of H1N1 influenza. Discussion meeting on Mathematical Models of Infection, Immunity and Inflammation, Indian Institute of Science, Bangalore. April 2017.

Invited Talks

- Data Science in Bioinformatics. Women in Data Science (WiDS) Mysuru, India, Sep 2020.
- Network Algorithms and their Applications in Biology. Sanjay Ghodawat University, Kolhapur, India, Dec 2018.
- Statistical Thinking in Biomedical Research. KLE College of Pharmacy, Bangalore, India, Apr 2018.

Graduate Coursework

- PhD (IISc): Current Trends in Drug Discovery, Special Topics in Theoretical Biology.
- MSc (NUS): Advanced Combinatorial Methods in Bioinformatics, Advanced Algorithms, Knowledge Discovery and Data Mining, Modeling and Analysis Techniques in Systems Biology, Advanced Topics in Data Mining.