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- Education and Experience
- Research Associate (Centre for Modeling and Simulation, Pune University)
  - Postdoctoral Fellow (NCBI, NIH)
  - Ph.D. in Computer Science (Duke University)
  - B.E. in Chemical Engineering (Pune University)
- Achievements
- Early Career Fellowship 2011, Wellcome-DBT
  - Fellows Award for Research Excellence, 2011 NIH
  - Ramanujan Fellowship 2010, DST Govt of India
- Research Subjects
- Machine Learning
  - Computational Biology
- Research Areas
- Genomics and Epigenomics
  - Transcriptional Regulation
  - Probabilistic Modeling
- Recent Publications
- L. Narlikar, et al. (2013) "One size does not fit all: On how Markov models order dictates performance of genomic sequence analyses". Nucleic Acids Research, 41:1416-24
  - L. Narlikar (2013) "MuMoD: a Bayesian approach to detect multiple modes of protein-DNA binding from genome-wide CHIP data". Nucleic Acids Research, 41:21-32
  - L. Narlikar and R. Jothi (2012) "ChIP-Seq data analysis: Identification of protein-DNA binding sites with SISSRs peak finder". Methods in Molecular Biology, 802:305-22
  - L. Narlikar et al. (2010) "Genome-wide discovery of human heart enhancers". Genome Research, 20:381-92