# Souvadra Hati

📞 +1-4709290024 | 💌 souvadrahati@gatech.edu | 🛅 souvadrahati

# **EDUCATION**

# **Georgia Institute of Technology**

2022 - Present

PhD in Computer Science • GPA: 4.0/4.0

Atlanta, GA

- Coursework: High Performance Computing, Advanced Computer Architecture
- Research Interests: Co-Design of Genomics and Graph Workloads, Asynchronous Parallel Algorithms

#### **Indian Institute of Science**

2018 - 2022

Bachelor of Science in Biology • GPA: 8.4/10.0

Bangalore, India

- Coursework: Computational Genomics, Machine Learning, Game Theory, Systems Biology
- Scholarship: Cargill Global Scholars fellow, KVPY scholar (All India Rank: 242)

## **PUBLICATIONS**

- [1] **S. Hati**, A. Hayashi, R. Vuduc, Asynchronous Distributed-Memory Parallel *k*-mer Counting, *IPDPS 2025*
- [2] S. Singhal\*, **S. Hati**\*, J. Young, V. Sarkar, A. Hayashi, R. Vuduc, Asynchronous Distributed-Memory Parallel Algorithms for Influence Maximization, *Supercomputing 2024*. <u>Link</u>
- [3] **S. Hati**, A. S. Duddu, M. K. Jolly, Operating Principles of Circular Toggle Polygons, *Physical Biology*, May 2021. Link
- [4] A. S. Duddu, S. Sahoo, **S. Hati**, S. Jhunjhunwala, M. K. Jolly, Multi-stability in cellular differentiation enabled by a network of three mutually repressing master regulators, *Journal of Royal Society Interface*, Sep 2020. Link

# **RESEARCH**

## **Asynchronous Genome Assembly**

2022 - Present

Advisor: Richard Vuduc

Georgia Tech

- Designed asynchronous k-mer counting algorithm, **2-10x** speedup over HySortK, **15-100x** speedup over KMC3.
- Developed parallel de-Bruijn graph construction algorithm; 2-5x speedup over PakMan.
- Implementation shows strong scaling up-to 6k CPU cores of Phoenix cluster at Georgia Tech.
- Preliminary work was selected as a finalist for the ACM SRC Poster competition at Supercomputing 2023.

#### **Influence Maximization on Social Network**

2023 - Present

Advisors: Richard Vuduc, Vivek Sarkar

Georgia Tech

- Designed two parallel algorithms for finding influencers on distributed social network graphs.
- First influence maximization software that enables weak scaling, and breaks the memory scalability barrier.
- Implementation shows 5-30x speedup over Ripples; strong scales up-to 8k CPU cores of NERSC Perlmutter.

#### **WORK EXPERIENCE**

#### Research Intern - Intel Labs

Summer 2024

Manager: Fabrizio Petrini

Intel Parallel Computing Labs

- Developed a multithreaded, AVX512-based software to simulate 3-hop interconnect networks.
- Reduced simulation time of large networks from 30-60 days to 2-6 hours.

# PAST PROJECTS

#### **Fast Genome Size Estimation**

2021 - 2022

Advisor: Chirag Jain Indian Institute of Science

- Developed MMC toolkit, a fast and multithreaded minimizer counter. [Link]
- Designed a parallel algorithm to sparsely sample DNA k-mers while covering all the nucleotides in the genome.
- Developed mathematical framework for *minimizer*-space genome size estimation.

# **Cell-Cell Interaction in Cancer Systems**

2020 - 2021

Advisor: Mohit Kumar Jolly

Indian Institute of Science

- Designed novel gene-regulatory network to model differentiation of CD4+ T-cells into different phenotypes.
- Developed MATLAB-based software to visualize the dynamics of biological networks.[Link]
- Best Poster at SMB 2021 conference for deciphering the operating principles of circular toggle polygon networks.

# **SKILLS**

**Languages:** C/C++, Python, MATLAB, R, Java, SQL, Julia **Libraries:** CUDA, MPI, OpenMP, OpenSHMEM, AVX

Tools: Git, LaTeX