Vivek Gopalakrishnan

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I am an first-year PhD student in Medical Engineering and Medical Physics at the Harvard-MIT Program in Health Sciences and Technology. My past research has focused on the development of computational methods to analyze multi-subject neuroimaging and cardiovascular data. Most recently, I developed statistical graph theory algorithms to perform biomarker discovery in network-valued maps of the brain. I also led a Design Team of undergraduate biomedical engineers to build a dynamic fusion image guidance system for minimally invasive heart surgery.

Education

8/21 – present Massachusetts Institute of Technology and Harvard Medical School, Cambridge, MA.

Doctor of Philosophy (Ph.D.) in Medical Engineering and Medical Physics.

- Advisor: Professor Polina Golland.
- o Concentration Area: Computer Science.
- 8/20 5/21 **Johns Hopkins University**, Baltimore, MD.

Master of Science and Engineering (M.S.E.) in Biomedical Engineering.

- Advisors: Professors Joshua T. Vogelstein (primary) and Carey E. Priebe (secondary).
- Concentration Area: Biomedical Data Science.
- Thesis: Multiscale Statistical Hypothesis Testing for k-Sample Graph Inference.
- 8/17 5/21 **Johns Hopkins University**, Baltimore, MD.

Bachelor of Science (B.S.) in Biomedical Engineering.

- Advisor: Professor René Vidal.
- Concentration Area: Biomedical Data Science.

Research and Engineering Experience

- 1/20 5/21 **Design Team Leader**, *Dept. of Biomedical Engineering*, Johns Hopkins University.
 - o Team leader and project manager for a team of eight undergraduate engineers.
 - Developed a multi-modal spatiotemporal registration algorithm to power a dynamic fusion image guidance system for minimally invasive heart surgery.
 - Skills used: Computer vision, image registration, cardiology, Python, and Git.
- 9/18 5/21 Undergraduate Research Assistant, Neurodata Lab, Johns Hopkins University.
 - Developed novel machine learning methods for analyzing populations of graph-valued objects [3, 1].
 - Discovered neuro-connectively similar subtypes of autism using joint embeddings of multi-network connectomes [4, 7].
 - Skills used: Graph theory, statistics, machine learning, Python, and Git.
- Summer 2018 **Summer Student Researcher**, *Ghebremichael Lab*, Ragon Institute of MGH, MIT and Harvard.
 - Developed predictive models of disease progression and immune recovery in HIV-positive children with TB co-infection [2].
 - Identified biomarkers for antiretroviral toxicity by applying machine learning methods to HIV-patient data [5].
 - Skills used: Survival analysis, ROC models, R, and Git.

- 12/17 5/18 **Design Team Member**, *Dept. of Biomedical Engineering*, Johns Hopkins University.
 - Designed and implemented a clinical machine learning algorithm to predict the onset of lung failure in pediatric patients.
 - **Skills used:** Representation learning, sparse dictionary learning, Python, and Git.
- 6/17 7/17 **Summer Researcher**, Fondazione Bruno Kessler, Trento, Italy.
 - Implemented a deep learning algorithm to quantitatively assess crop damage from images taken by farmers.
 - Skills used: Computer vision, Keras, Python, and Git.
- 6/16 6/17 **High School Researcher**, *Tufts University BioSeq Program*, Tufts University.
 - Used next-generation sequencing (NGS) to generate a novel human oral microbiome data set.
 - Analyzed sequence data using dimensionality reduction and hidden Markov models to identify microbial biomarkers of oral health.
 - o Presented at the 2017 Intel International Science and Engineering Fair, Los Angeles, CA.
 - **Skills used:** DNA library construction, next-generation sequencing, dimensionality reduction, hidden Markov models, and R.

Publications and Preprints

- [1] Jaewon Chung, Eric Bridgeford, Jesus Arroyo, Benjamin D. Pedigo, Ali Saad-Eldin, **Vivek Gopalakrishnan**, Liang Xiang, Carey E Priebe, and Joshua T Vogelstein. "Statistical Connectomics". *Annual Review of Statistics and Its Application* (Mar. 2021). DOI: https://doi.org/10.1146/annurev-statistics-042720-023234.
- [2] Vivek Gopalakrishnan, Eliezer Bose, Usha Nair, Yuwei Cheng, and Musie Ghebremichael. "Pre-HAART CD4+ T-Lymphocytes as Biomarkers of Post-HAART Immune Recovery in HIV-Infected Children with or without TB Co-Infection". BMC Infectious Diseases (Oct. 2020). DOI: 10.1186/s12879-020-05458-w.
- [3] **Vivek Gopalakrishnan**, Jaewon Chung, Eric Bridgeford, Benjamin D. Pedigo, Jesús Arroyo, Lucy Upchurch, G. Allan Johnson, Nian Wang, Youngser Park, Carey E. Priebe, and Joshua T. Vogelstein. "Multiscale Comparative Connectomics". *arXiv:2011.14990* (Nov. 2020). arXiv: 2011.14990.
- [4] Nian Wang, Robert J Anderson, David G Ashbrook, **Vivek Gopalakrishnan**, Youngser Park, Carey E Priebe, Yi Qi, Rick Laoprasert, Joshua T Vogelstein, Robert W Williams, and G Allan Johnson. "Variability and Heritability of Mouse Brain Structure: Microscopic MRI Atlases and Connectomes for Diverse Strains". *NeuroImage* (Aug. 2020). DOI: 10.1016/j.neuroimage.2020.117274.
- [5] Jong Soo Lee, Elijah Paintsil, Vivek Gopalakrishnan, and Musie Ghebremichael. "A Comparison of Machine Learning Techniques for Classification of HIV Patients with Antiretroviral Therapy-Induced Mitochondrial Toxicity from Those Without Mitochondrial Toxicity". BMC Medical Research Methodology (Nov. 2019). DOI: 10.1186/s12874-019-0848-z.

Conference Presentations

- [6] Vivek Gopalakrishnan and Joshua T Vogelstein. "Statistical Methods for Multiscale Comparative Connectomics". NeuroMatch. Oct. 2020. URL: https://www.neuromatch.io/abstract?submission_id=recARY71P6SjXY3xK.
- [7] Vivek Gopalakrishnan and Joshua T Vogelstein. "Towards Discovering Heterogeneity in Autism via Multi-Network Connectomics". Biomedical Engineering Society (BMES). Philadelphia, PA, Oct. 2019.

Skills

Programming Python, R, Julia, git, LATEX

- DevOps GitHub, AWS, Docker
 - APIs Python Data Science Stack, Tensorflow, PyTorch
- Bioinformatics Microbiome and B-/T-cell repertoire analysis
 - Wet Lab Microfabrication, 3-D tissue culture, NGS library construction

Teaching

- Spring '21 **Head TA NeuroData Design II**, *EN.BME.438/638*, Johns Hopkins University.
 - Fall '20 **Head TA NeuroData Design I**, *EN.BME.437/637*, Johns Hopkins University.
 - Fall '20 **TA Linear Algebra**, *AS.MATH.201*, Johns Hopkins University.
 - Fall '20 **TA Computational Cardiology Lab**, *EN.BME.487*, Johns Hopkins University.
 - Fall '18 Head PILOT Leader, Dept. of Academic Support, Johns Hopkins University.
- to Spring '20 \circ Led group tutoring sessions (≈ 10 students/group), and wrote weekly problem sets and lectures.
 - Led weekly meetings to train junior PILOT leaders in mathematics and pedagogy.

Fellowships and Awards

- 2020 **Provost's Undergraduate Research Award**, *Office of Undergraduate Research*, Johns Hopkins University.
- 2019 INBT Research Award, Institute for NanoBioTechnology, Johns Hopkins University.
- 2019 **Joseph C. Pistritto Research Fellowship**, *Dept. of Computer Science*, Johns Hopkins University.
- 2018 **AWS Cloud Credits for Research Grant**, *Dept. of Computer Science*, Johns Hopkins University.
- 2017 **Second Place Winner**, *Intel International Science and Engineering Fair (ISEF)*, Category: Microbiology.
- 2017 Internship Award, Intel ISEF Special Award, Fondazione Bruno Kessler.
- 2017 **Semi-Finalist**, Regeneron Science Talent Search, Microbiology.
- 2017 **Second Place Winner**, Massachusetts State Science & Engineering Fair.
- 2016 Second Place Winner, Massachusetts State Science & Engineering Fair.