Kasthuri Kannan, PhD

Associate Professor, UT MD Anderson Cancer Center

≤ kskannan@mdanderson.org 🛛 📞 (551) 284 9106

06 • 2130 W Holcombe Blvd. Houston. TX 77494

in linkedin.com/in/kasthuri-kannan-0a05b43

PROFILE

As a seasoned data scientist with 15 years in life sciences, I excel in managing complex, large-scale data analysis and modeling projects. I am recognized for a strong leadership, guiding a multi-institutional team of 6 scholars and 7 faculty members. Actively seeking to leverage my expertise in an industrial setting, where I can drive innovation and deliver impactful results.

EDUCATION

Doctor of Philosophy (Computer Science) Texas A&M University

Master of Science (Mathematics) Texas A&M University

Master of Science (Mathematics) Indian Institute of Technology, Madras

Bachelor of Science (Mathematics) University of Madras

SKILLS

Leadership ● ● ● ● ● ● ● ■ Managing data science and bioinformatics projects

Software (Representative) • • • • • • • • Python, R, Java, Cypher, SQL, Bioinformatics tools, Unix/HPC/Bash, Neo4j, MySQL, PostgreSQL, HTML, Javascript, Spatstat, Pytorch, Tensorflow

PROFESSIONAL EXPERIENCE

UT MD Anderson Cancer Center

Associate Professor

- Leading graph database development for multi-omics data integration
- Establishing spatial modeling of pathology images using R spatstat
- Supervising post-doctoral fellows and students

New York University

Assistant Professor

- Established bioinformatics pipelines and investigated cancer datasets
- Directed data science courses see Data Science Education section

Memorial Sloan-Kettering Cancer Center

Research Fellow/Associate

- Established mutation pipeline for Brain/Head & Neck cancers
- Discovered ATRX mutations in lower grade gliomas
- Provided directed insights in various cancer studies through data analysis

2002 – 2008 College Station, Texas

2000 – 2002 College Station, Texas

> 1998 – 2000 Chennai, India

1995 – 1998 Chennai, India

Data Science/Machine Learning/Al • • • • • Math, CS, Statistics and Programming Ethical hacking, Domain knowledge, LLMs

Bioinformatics & Image Processing • • • • • • Pipeline development and data engineering Geospatial analysis

02/2020 – present Houston, TX

11/2013 – 08/2019 New York, NY

04/2011 – 10/2013 New York, NY

Pennsylvania State University

Research Associate

- Organized data analysis pipelines
- Offered bioinformatics consultation to researchers
- Managed sequencing tasks

Stowers Institute for Medical Research

Research Specialist II

- Developed image processing methods for worm and fly tracking
- Implemented an automated data workflow to process yeast cell images
- Responsible for high quality image acquistion and microscope maintanence

Knowledge Based Systems, Inc

Internship

- Delivered data driven directed insights for aircraft movement operations
- Proposed cost and time saving measures for managing air force logistics

ACTIVE DATA SCIENCE PROJECTS

Geospatial modeling of tissues — Spatial point processes are powerful statistical frameworks for studying point patterns. By representing cells as points and annotating the measurements taken on those cells, such as gene expression at single cell level, it is appropriate to study their interactions using point processes. We extensively use spatstat, an R package, to model interactions and derive directed insights in cancers.

Biomarker discovery using graph database — Graph databases can help identify biomarkers by efficiently representing complex biological networks. By enabling powerful queries and community detection algorithms, graph databases make exploring the relationships between multiple genes easier, thus facilitating the discovery of potential biomarkers critical for diagnostics or therapeutic targets. We use Neo4j, a property graph database, and algorithms from the Graph Data Science Library (GDS) to derive insights and propose actionable biological targets.

Biomarker validation using Graph Neural Networks (GNNs) — Insilico validation of biomarkers is critical for providing actionable biological targets for experiments and prognosis. We apply GNNs (convolutional, attention etc.) to validate critical biomarkers discovered through the graph database. GNNs enhance biomarker discovery by leveraging the structure of biological networks, such as genegene interactions, and learning meaningful node/edge representations. By aggregating information from a node's neighbors, GNNs capture local and global patterns, allowing the model to predict relationships between genes and diseases more effectively.

DATA SCIENCE EDUCATION

Developed and taught the following data science courses and lectures

- Programming for Data Analysis
- Machine Learning and Al
- Methods in Quantitative Biology

INVITED TALKS (REPRESENTATIVE)

Mayo Clinic (2023), National University of Singapore (2022), Texas A&M University (2021), Courant Institute of Mathematical Sciences, w/ honorarium (2018)

PUBLICATIONS

Authored/co-authored 32 peer reviewed articles, that includes very high-impact journals. Please refer https://pubmed.ncbi.nlm.nih.gov/?term=kasthuri+kannan&sort=date&sort_order=desc 🛛

01/2008 – 09/2010 Kansas City, MO

01/2007 – 12/2008 College Station, TX