

R RANGA SUDHARSHAN

rsudhars@med.umich.edu ◊ Personal Website

EDUCATION

Indian Institute of Technology (BHU), Varanasi

Oct 2023

Integrated Dual Degree (B.Tech + M.Tech) in Bioengineering and Biomedical Technology

CGPA: 8.63/10.0

WORK EXPERIENCE

University of Michigan, Ann Arbor

Nov 2023 - Present

Lead Bioinformatician, Pancreatic Cancer Lab

[link]

Research Specialist Lab Associate, Systems Imaging and Bioinformatics Lab

[link]

Visa: H1B (Valid till Nov 2026)

JOURNAL PAPER PREPRINTS

TIGIT as a mediator of immune suppression in COMPASS-like complex gene-mutated pancreatic cancer S. Zhang, E. R. Daniels, J.M. Ranga Sudharshan, H. C. Kim, D.G.T, S. Krishnan, T. L. Frankel, E. Hissong, A. Rao, N. Assarzadegan, J. Shi *Accepted - Modern Pathology (I.F: 5.5)*

QHGM: Quantum Hamiltonian-Based Generative Model for Gene Regulatory Network Inference
M. Sohail, Ranga Sudharshan, S. Pradhan, A. Rao
In Review - Nature Methods

Rewiring Oncogenic Transcriptional Complexes with Domain-ALTeration Chimeras (DALTACs) in Prostate Cancer J. Luo, J. Yang, J. C. Tien, ... Ranga Sudharshan, A. Rao, A. Chinnaiyan, et. al
In Review - Nature Genetics

CONFERENCE PAPERS AND ABSTRACTS

QubitLens: An Interactive Learning Tool for Quantum State Tomography

[link]

M.A.Sohail, Ranga Sudharshan S. Pradhan, A. Rao [QSEEC IEEE Quantum Week 2025, ABQ]

Spatial multi-omics reveal distinct immunosuppressive lipid-laden macrophages in primary CNS lymphoma compared to systemic DLBCL [Co- author (ASH 2025, FL – oral presentation)]

[link]

Abstract: TIGIT-mediated immune suppression in KMT2D-mutant pancreatic cancer

[Co author (AACR 2025)]

[link]

Abstract: Epigenetic Tumor Suppressor KMT2D Shapes the Immune Landscape in Pancreatic Cancer [Co-author (AACR 2025)]

[link]

Abstract: KMT2D Loss Induces Enrichment of Plod2+ Fibroblasts in Pancreatic Cancer

[Co-author (AACR 2025)]

[link]

Comparing Perturbagens from Differential Gene Expression Data Analysis of ASD using Random Forest and Statistical Test. [(Current Directions in Biomedical Engineering, 2023)]

Ranga Sudharshan, S. K., P., B.M.G., R. Jac.F.A.

[link]

AWARDS, TALKS AND OUTREACH

DAAD (KOSPIE) Scholar – Awarded Combined Study and Practice Stays Scholarship (declined).

IBGAA Graduate Student Admission Scholarship, IIT BHU Alumni Association (\$700).

Gave a talk titled "Simplifying quantum state tomography using maximum likelihood estimation"
IEEE International Symposium on Information Theory 2025 [\[link\]](#)

Founding Member, Research Community, IIT BHU
Mentored 25+ sophomores for research internships. [\[link\]](#)

ONGOING RESEARCH PROJECTS

Spatial Analysis of Dormant Breast Cancer Cells in the Bone Marrow Microenvironment

Collaborator: Dr. Grace Bushnell (University of Minnesota)

- Processed 2 μ m Visium HD spatial transcriptomics data using Bin2Cell for segmentation and deconvoluted with a bone marrow single-cell atlas via RCTD; registered cytokeratin-positive cells from IHC and annotated tumor cells with nearest-neighbor mapping.
- Performed neighborhood enrichment and computed Moran's I to assess spatial co-localization and auto-correlation of cell types and top differentially expressed genes.
- Applied GMM clustering to identify tumor cell clusters and compared the immune microenvironment of clustered versus singlet tumor cells.

Transcriptomic Profiling of Alveolar Type II Cell Memory Programs using scRNA-seq

Collaborator: Dr. Anukul Shenoy (Department of Microbiology and Immunology, UM Ann Arbor)

- Performed scRNA-seq analyses on FACS-purified AT2 cells, validating conserved interferon-induced (MHC-I) and adaptive (MHC-II) memory programs.
- Conducted differential expression and pathway enrichment analyses (KEGG, GO, MSigDB) using clusterProfiler and inferred transcription factor activity with Decoupler (ULM + CollecTRI database) to characterize regulatory programs.
- Integrated microarray datasets, consolidating diverse perturbation phenotypes (STAT1-, IL-17A/F-, IFN- γ) to strengthen mechanistic models of epithelial memory.

Variational Hamiltonian Learning of Cell-Cell Networks Across Tumor Progression Landscapes

Collaborator: Dr. Sandeep Pradhan (Dept. of Electrical Engineering, UM Ann Arbor)

- Converted CODEx image patches into continuous cell-density trajectories using kernel density estimation and ordered tumor cells into pseudo-temporal “quantiles” to approximate progression.
- Applied variational Hamiltonian learning using an Ising model, where cell-density trajectories serve as time-dependent observations to recover interaction weights of a parameterized Hamiltonian.
- Engineered POVM-based likelihood estimation with tensor contractions for 4^n outcome probabilities and fully JIT-compiled optimization using JAX and Optax.

TEACHING EXPERIENCE

Teaching Assistant Jan 2023 – May 2023
Computational Neuroscience Lab, School of Biomedical Engineering, IIT (BHU)

Teaching Assistant Aug 2022 – Nov 2022
Nano-Cellular Medicine and Biophysics Lab, School of Biomedical Engineering, IIT (BHU)

TECHNICAL SKILLS

Programming	Python, R, C; Git; Bash (Slurm HPC)
Quantum/Computational Tools	PennyLane, Qiskit, TensorFlow, PyTorch, scikit-learn, NLTK
Bioinformatics Tools	Scanpy, scVI, Seurat, ArchR, C-Side, Decoupler, InferCNV, Giotto