

R RANGA SUDHARSHAN

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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 *In Review - 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
Submitted to 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
Submitted to 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