# Jai Prakash Veerla

Ph.D. Candidate, The University of Texas at Arlington SEIR Buildingn Room 428F 655 W Mitchell Street Arlington, TX 76010 ☑ jxv6663@mavs.uta.edu ﴿ jaiprakashveerla.com

### Education

2022-Present **Ph.D. in Computer Science**, *University of Texas at Arlington*, TX

Computer Science & Engineering @ UTA, GPA:- 3.91/4.0

Advisor: Dr. Jacob Luber

Dissertation: "Enhancing Cancer Imaging: Spatial Visualization, 3D Pathology, Model

Trustworthiness, with Immersive Technology"

2021–2022 **M.S. in Computer Science**, *University of Texas at Arlington*, TX

Computer Science & Engineering @ UTA, Converted to Ph.D.

2016-2020 B.Tech. in Computer Science & Engineering First Class with Distinction, INTUH, Hyderabad, IN

# Relevant Experiences

Nov 2021 - Graduate Research Assistant, The University of Texas at Arlington

Present Arlington, TX, 76010, Multi-Interprofessional Center for Health Informatics (MICHI)

Oct 2020 - Data Analyst, Vamstar, Hyderabad, IN

Feb 2021

Jan 2020 - Research Scientist Intern, Defence Research and Development Organisation (DRDO),

Sept 2020 Hyderabad, IN

RCMA (Missiles)

Sept 2019 - Machine Learning Research Intern, Risk Edge Solutions, Hyderabad, IN

Jan 2020

May 2019 - Research Intern, The International Institute of Information Technology Hyderabad

Aug 2019 (IIIT Hyderabad), Hyderabad, IN

Language Technologies Research Center (LTRC)

### Proficiencies

#### **Languages and Technologies**

Python (numpy, pandas, scipy, matplotlib, sklearn, pytorch, pytorch lightning), Machine Learning libraries, Deep Learning architectures, Computer Vision, MLOPs, C++, CUDA, CoreML, MLX Framework, Xcode, Swift, R (ggplot, shiny), bash, MySQL, ŁTĘX, Git, Linux, HPC (slurm), Abode Creative Cloud (Illustrator, Photoshop, Premiere Pro, etc.), HTML5, CSS, Docker, Kubernetes, Unity, Tableau, AWS (EC2, load balancing, etc.)

# Publications

(\* denotes co-first authors)

- 1 **Jai Prakash Veerla**, Partha Sai Guttikonda, Amir Hajighasemi, Jillur Rahman Saurav, Aarti Darji, Cody T Reynolds, Mohamed Mohamed, Mohammad S Nasr, Helen H Shang, and Jacob M Luber. Spatialvisvr: An immersive, multiplexed medical image viewer with contextual similar-patient search. *arXiv preprint arXiv:2401.02882*, 2024.
- 2 **Jai Prakash Veerla\***, Poojitha Thota\*, Partha Sai Guttikonda, Shirin Nilizadeh, and Jacob M Luber. Vulnerabilities unveiled: Adversarially attacking a multimodal vision language model for pathology imaging. *arXiv preprint arXiv:2401.02565*, 2024.
- 3 **Jai Prakash Veerla**, Jillur Rahman Saurav, Michael Robben, and Jacob M Luber. Analyzing lack of concordance between the proteome and transcriptome in paired scrna-seq and multiplexed spatial proteomics. *arXiv preprint arXiv:2307.00635*, 2023.
- 4 Helen H Shang\*, Mohammad Sadegh Nasr\*, **Jai Prakash Veerla**, Parisa Boodaghi Malidarreh, MD Saurav, Amir Hajighasemi, Manfred Huber, Chace Moleta, Jitin Makker, and Jacob M Luber. Histopathology slide indexing and search: Are we there yet? *arXiv preprint arXiv:2306.17019*, 2023.
- 5 Michael Robben, Amir Hajighasemi, Mohammad Sadegh Nasr, **Jai Prakash Veerla**, Anne M Alsup, Biraaj Rout, Helen H Shang, Kelli Fowlds, Parisa Boodaghi Malidarreh, Paul Koomey, et al. The state of applying artificial intelligence to tissue imaging for cancer research and early detection. *arXiv preprint arXiv:2306.16989*, 2023.
- 6 Michael Robben, Mohammad Sadegh Nasr, Avishek Das, **Jai Prakash Veerla**, Manfred Huber, Justyn Jaworski, Jon Weidanz, and Jacob Luber. Comparison of the strengths and weaknesses of machine learning algorithms and feature selection on kegg database microbial gene pathway annotation and its effects on reconstructed network topology. *Journal of Computational Biology*, 30(7):766–782, 2023.
- 7 Amir Hajighasemi\*, MD Saurav\*, Mohammad S Nasr, **Jai Prakash Veerla**, Aarti Darji, Parisa Boodaghi Malidarreh, Michael Robben, Helen H Shang, and Jacob M Luber. Multimodal pathology image search between h&e slides and multiplexed immunofluorescent images. *arXiv preprint arXiv:2306.06780*, 2023.
- 8 Neel R Vora\*, Amir Hajighasemi\*, Cody T Reynolds, Amirmohammad Radmehr, Mohamed Mohamed, Jillur Rahman Saurav, Abdul Aziz, **Jai Prakash Veerla**, Mohammad S Nasr, Hayden Lotspeich, et al. Real-time diagnostic integrity meets efficiency: A novel platform-agnostic architecture for physiological signal compression. *arXiv* preprint arXiv:2312.12587, 2023.
- 9 Parisa Boodaghi Malidarreh\*, Biraaj Rout\*, Mohammad Sadegh Nasr\*, Priyanshi Borad\*, Jillur Rahman Saurav\*, **Jai Prakash Veerla**, Kelli Fenelon, Theodora Koromila, and Jacob M Luber. Predicting future states with spatial point processes in single molecule resolution spatial transcriptomics. *arXiv preprint arXiv:2401.02564*, 2024.

# Teaching

**TA Spring 2024 @ UTA Department of Computer Science**, *College of Engineering* Bioinformatics (CSE 5370)

**TA Fall 2021 @ UTA Department of Physics**, *College of Science* Python for Data Science (DATA 3402)

### Honors and Awards

- Fall 2021 Lonestar Scholarship, University of Texas at Arlington
- Fall 2021 Silverstar Scholarship, University of Texas at Arlington
- Fall 2020 First Class with Distinction, JNTUH

### Social Media and Code

LinkedIn, https://www.linkedin.com/in/jai-prakash-veerla/

Google Scholar, https://scholar.google.com/citations?user=0EHcHsgAAAAJhl=en

Github, https://github.com/jaiprakash1824

Twitter, https://twitter.com/jp\_veerla

### Relevant Coursework

### Completed @ University of Texas at Arlington

Special Topics in Advanced Multimedia Graphics, & Image Processing, Special Topics in Advanced Computer Science, Machine Learning, Bioinformatics, Artificial Intelligence, Numerical Methods, Design and Analysis of Algorithms, Data Analysis & Modeling Techniques, Computer Architecture, Distributed Systems, Computer Vision (Audited), Neural Networks (Audited)

# Conference Presentations, Posters, and Invited Talks

1 Jai Prakash Veerla, Jillur Rahman Saurav, Michael Robben, and Jacob M. Luber. Analyzing lack of concordance between the proteome and transcriptome in paired scrna-seq and multiplexed spatial proteomics [short paper talk]. *IEEE CIBCB 2023*, August 2023. Eindhoven, NL.