

|                     |   |  |
|---------------------|---|--|
| CONTACT INFORMATION | linkedin.com/in/aarthi-venkat/<br>aarthivenkat.github.io  | (408) 799-9189<br>avenkat@broadinstitute.org   |
| EDUCATION           | <b>Yale University</b><br><b>Ph.D.</b> in Computational Biology & Bioinformatics<br><b>M.S.</b> in Computational Biology & Bioinformatics<br><br><b>The University of California, San Diego</b><br><b>B.S.</b> in Bioengineering: Bioinformatics  | May 2024<br>Dec 2021<br><br>Jun 2019   |
| RESEARCH EXPERIENCE | <b>Eric and Wendy Schmidt Center Postdoctoral Fellow</b><br>Broad Institute of MIT & Harvard, Drs. Marinka Zitnik & Nir Hacohen <ul style="list-style-type: none"><li>Bridging systems biology with geometric deep learning and foundation models toward transferring insights across biological scales and contexts</li></ul><br><b>Computational Biology &amp; Bioinformatics Ph.D. Student</b><br>Yale University, Dr. Smita Krishnaswamy <ul style="list-style-type: none"><li>Developed geometric representation learning methods to characterize cellular and molecular behavior in diverse, co-led collaborations</li></ul><br><b>Applied Science Research Intern</b><br>Google Brain, Drs. Lucy Colwell & Farhad Hormozdiari <ul style="list-style-type: none"><li>Performed ML-guided biological sequence design with Google Genomics, UCSF</li><li>Achieved top performance for CRISPR RNA guide efficacy and expression prediction</li></ul><br><b>Bioinformatics Research Assistant</b><br>La Jolla Institute for Immunology, Dr. Ferhat Ay <ul style="list-style-type: none"><li>Characterized 3D structure of malaria-related parasite genomes from Hi-C sequencing</li><li>Corrected <i>Toxoplasma gondii</i> misassembly with Hi-C and long-read sequencing</li></ul><br><b>Computational Biology Research Assistant</b><br>Institute for Genomic Medicine, Dr. Theresa Gaasterland <ul style="list-style-type: none"><li>Performed bioinformatic analysis of primary congenital glaucoma exomes</li></ul><br><b>Genome Informatics Intern</b><br>Regeneron Pharmaceuticals, Regeneron Genetics Center <ul style="list-style-type: none"><li>Integrated loss-of-function variant and target annotation for over 500,000 exomes</li><li>One of 7 selected out of 250+ interns to present at company-wide annual event</li></ul><br><b>Data Analytics Intern</b><br>Auris Health, Research & Development <ul style="list-style-type: none"><li>Built cloud-based pipeline to facilitate high-performance analysis of endoscopic robot</li></ul> | Sept 2024-Present<br><br>Aug 2019-Jul 2024<br><br>Sept 2021-Dec 2021<br><br>Oct 2016-Sept 2019<br><br>Sept 2018-Aug 2019<br><br>Jun 2018-Sept 2018<br><br>Jun 2017-Sept 2017 |
| PUBLICATIONS        | *§ Denote equal contribution.<br>Links to full publications available on my website: <a href="https://aarthivenkat.github.io">https://aarthivenkat.github.io</a><br><br>[1] C Garcia*, <b>A Venkat*</b> , DC McQuaid* ... S Krishnaswamy <sup>§</sup> , MD Muzumdar <sup>§</sup> . <i>Beta cells are essential drivers of pancreatic ductal adenocarcinoma development</i> . In Revision at Nature Communications.<br>[2] <b>A Venkat*</b> , S Youlten*, BP San Juan* ... S Krishnaswamy <sup>§</sup> , CL Chaffer <sup>§</sup> . <i>AAnet resolves a continuum of spatially-localized cell states to unveil tumor complexity</i> . Cancer Discovery (2025).<br>[3] <b>A Venkat</b> , S Leone, S Youlten, E Fagerberg, J Attanasio, NS Joshi, S Krishnaswamy. <i>Mapping the gene space at single-cell resolution with gene signal pattern analysis</i> . Nature Computational Science (2024).  |  |

- [4] **A Venkat\***, J Chew\*, F Cardoso Rodriguez, CJ Tape, M Perlmuter<sup>§</sup>, S Krishnaswamy<sup>§</sup>. *Directed scattering for knowledge graph-based cellular signaling analysis*. ICASSP (2024).
- [5] **A Venkat\***, M Carlino\*, B Lawton\* ... S Krishnaswamy<sup>§</sup>, D Krause<sup>§</sup>. *Single-cell analysis reveals transcriptional dynamics in primary parathyroid tissue*. Genome Research (2024).
- [6] **A Venkat**, D Bhaskar, S Krishnaswamy. *Multiscale geometric and topological analyses for characterizing and predicting immune responses from single-cell data*. Cell Trends in Immunology (2023).
- [7] D Bhaskar\*, DS Magruder\*, M Morales, E De Brouwer, **A Venkat**, F Wenkel, J Noonan, G Wolf, N Ivanova, S Krishnaswamy. *Inferring dynamic regulatory interaction graphs from time series data with perturbations*. LoG Conference (2023).
- [8] S Leone, A Tong, G Huguet, **A Venkat**, G Wolf, S Krishnaswamy. *Graph Fourier MMD for Signals on Graphs*. SampTA (2023).
- [9] X Sun\*, S Gupta\*, A Tong\*, M Kuchroo\*, D Bhaskar\*, C Liu, **A Venkat** ... CL Chaffer<sup>§</sup>, S Krishnaswamy<sup>§</sup>. *Revealing dynamic temporal trajectories and underlying regulatory networks with Cflows*. In Review.
- [10] M Damo, N Hornick, **A Venkat** ... NS Joshi. *PD-1 prevents pathogenicity of effector CD8 T cells that infiltrate skin under homeostatic conditions*. Nature (2023).
- [11] M Amodio, SE Youtlen, **A Venkat**, BP San Juan, CL Chaffer, S Krishnaswamy. *Single-cell multi-modal GAN reveals spatial patterns in single-cell data from triple-negative breast cancer*. Cell Patterns (2022).
- [12] KA Connolly, M Kuchroo, **A Venkat** ... NS Joshi. *A reservoir of stem-like CD8+ T cells in the tumor draining lymph node preserves the ongoing antitumor immune response*. Science Immunology (2021).
- [13] Y Su\*, **A Venkat\***, Y Yadav, L Puglisi, S Fodeh. *Twitter-based analysis reveals differential COVID-19 concerns across areas with socioeconomic disparities*. CBM (2021).
- [14] J Xia, **A Venkat**, ML Reese, KG Le Roch, F Ay, JP Boyle. *Third generation sequencing revises the molecular karyotype for Toxoplasma gondii and identifies emerging copy number variants in sexual recombinants*. Genome Research (2021).
- [15] EM Bunnik, **A Venkat\***, J Shao\*, KE McGovern ... F Ay<sup>§</sup>, KG Le Roch<sup>§</sup>. *Comparative 3D Organization in Apicomplexan Parasites*. PNAS (2019).

## PRESENTATIONS

- *Analysis of Therapeutic Response in Hepatocellular Carcinoma with Graph-based Machine Learning Methods on Spatial Transcriptomics Data*  
ASCB Poster Presentation (2025)
- *Mapping the gene space at single-cell resolution with gene signal pattern analysis*  
LOG Poster Presentation (2025)
- *Epistemic Responsibility and Interdisciplinarity in the Age of AI for Science*  
LLMs and Digital Autonomy: From Misinformation to Context Collapse Oral Presentation (2025)
- *Dissecting cellular and molecular mechanisms of pancreatic cancer with deep learning*  
ISMB MLCSB Conference Oral Presentation (2025)
- *Dissecting cellular and molecular mechanisms of pancreatic cancer*  
Dr. Samuel M. Nabrit Conference, Brown University, Oral Presentation (2025)
- *Geometric representation learning for single-cell biology across contexts*  
Invited talk, Brown University (2025)
- *Learning cellular and molecular mechanisms of pancreatic cancer*  
Connecting the Dots Broad Institute Symposium Poster Presentation (2024)
- *Mapping the gene space at single-cell resolution with gene signal pattern analysis*  
Yale Department of Genetics Symposium Poster Presentation (2023)
- *Mapping the gene space at single-cell resolution with gene signal pattern analysis*  
Gruber Science Fellowship Symposium Poster Presentation (2023)

- *Learning directed and hyperbolic embeddings*  
Graph Signal Processing Workshop Oral Presentation (2023)
- *PHATE reveals cell state transformation in Tercen biomedical data analysis platform*  
CYTO Oral Presentation (2023)
- *Mapping the gene space at single-cell resolution with gene signal pattern analysis*  
AAI Immunology Poster Presentation (2023)
- *Elucidating mechanisms of endocrine-exocrine signaling in pancreatic cancer*  
Yale Single Cell Symposium Oral Presentation (2022)
- *Manifold-based gene density estimates reveal immune signaling in meningioma*  
ISMB MLCSB Conference Poster Presentation (2021)
- *Archetypal analysis of antigen-specific T cell responses across conditions*  
CSHL Systems Immunology Conference Poster Presentation (2021)
- *Leveraging the Power of Human Genetics through Knockout Discovery*  
Regeneron Oral Presentation & Poster (2018)

|                           |  |                        |
|---------------------------|--|------------------------|
| TEACHING<br>EXPERIENCE    | <b>Teaching Assistant, Computational Genomics</b><br>Cold Springs Harbor Laboratory Workshop   | Nov 2022, Dec 2023     |
|                           | <ul style="list-style-type: none"> <li>• Designed and presented single-cell workshops for 20-40 PhD-level researchers</li> </ul>   |                        |
|                           | <b>Teaching Fellow, Deep Learning Theory and Applications</b><br>Yale University, Computer Science   | S 2021, S 2024         |
|                           | <ul style="list-style-type: none"> <li>• Held recitations, designed and graded homework, exams, and projects for undergraduate and graduate students</li> </ul>  |                        |
|                           | <b>Teaching Assistant, Machine Learning for Single-cell Analysis</b><br>Yale University, Department of Genetics & Yale SEAS  | May 2020, Jan 2021     |
| FELLOWSHIPS<br>AND GRANTS | <ul style="list-style-type: none"> <li>• Co-taught 100+ researchers across all levels in tools for single-cell analysis</li> </ul>   |                        |
|                           | <b>Teaching Assistant, Introduction to Biomedical Data Science and Health Informatics</b><br>Yale Center for Medical Informatics   | Jun 2020               |
|                           | <ul style="list-style-type: none"> <li>• Assisted in Python for biomedical data analysis for researchers across all levels</li> </ul>  |                        |
|                           | <b>Genetics Undergraduate Tutor / Instructional Assistant</b><br>UC San Diego Biological Sciences  | F 2017, F 2018, S 2019 |
|                           | <ul style="list-style-type: none"> <li>• Developed material for weekly recitation sessions, office hours, and exam preparation</li> <li>• Received Excellence in Teaching Award for top performance (100% positive reviews)</li> </ul> |                        |
| FELLOWSHIPS<br>AND GRANTS | <b>GSA Conference Travel Fellowship</b><br>Yale University   | Jun 2023, Mar 2024     |
|                           | <b>Yale Gruber Science Fellowship</b><br>Yale University   | Aug 2019               |
| HONORS AND<br>AWARDS      | <ul style="list-style-type: none"> <li>• Most prestigious award offered by Graduate School of Arts and Sciences to incoming science PhDs in recognition of outstanding accomplishments and promise</li> </ul>                          |                        |
|                           | <b>Public Communication Certificate</b><br>Poorvu Center for Teaching & Learning, Yale University  | 2023                   |
|                           | <ul style="list-style-type: none"> <li>• Certificate for skills developed in oral and written communication</li> </ul>   |                        |
|                           | <b>OHHER Award Finalist for Yale Research Excellence</b><br>Yale School of Medicine, Office of Health Equity Research  | 2022                   |
|                           | <ul style="list-style-type: none"> <li>• Received for “Twitter-based analysis reveals differential COVID-19 concerns across areas with socioeconomic disparities”</li> </ul>   |                        |
| HONORS AND<br>AWARDS      | <b>Outstanding Academic Achievement in Bioengineering</b><br>The University of California, San Diego   | 2019                   |
|                           | <ul style="list-style-type: none"> <li>• Highest performance in graduating class in Bioengineering: Bioinformatics</li> </ul>  |                        |

|                     |  |            |
|---------------------|--|------------|
|                     | <b>Excellence in Teaching Award</b><br>The University of California, San Diego   | 2019       |
|                     | <ul style="list-style-type: none"> <li>Highest performance evaluation for teaching assistance in Genetics</li> </ul>   |            |
|                     | <b>Tau Beta Pi Engineering Honors</b><br>The University of California, San Diego   | 2018, 2019 |
|                     | <ul style="list-style-type: none"> <li>Awarded to engineering students displaying high academic achievement and personal, professional integrity</li> </ul>  |            |
|                     | <b>Muir College Caledonian Honors</b><br>The University of California, San Diego   | 2018, 2019 |
|                     | <ul style="list-style-type: none"> <li>Awarded to engineering students displaying high academic achievement and personal, professional integrity</li> </ul>  |            |
|                     | <b>Provost Honors</b><br>The University of California, San Diego   | 2015-2019  |
|                     | <ul style="list-style-type: none"> <li>Received 12 times for high academic achievement</li> </ul>  |            |
| ACADEMIC<br>SERVICE | <b>Research Mentor</b> for Sofia Lara (MIT Biological Engineering)   | 2025       |
|                     | <b>Research Mentor</b> for Hannah Thomas (Nashua High School South)  | 2025       |
|                     | <b>Invited Reviewer</b> Computational & Structural Biotechnology Journal   | 2025       |
|                     | <b>Broad Summer Research Program</b> Selection Committee   | 2025       |
|                     | <b>Program Committee Member</b> for ISMB/ECCB 2025   | 2025       |
|                     | <b>Foundation Models in Genomics Panel Moderator</b> @ Broad   | 2024       |
|                     | <b>Invited Reviewer</b> RECOMB 2024  | 2024       |
|                     | <b>Invited Reviewer</b> Yale Journal of Biology and Medicine   | 2023       |
|                     | <b>Student Advisory Board</b> , Poorvu Center for Teaching & Learning  | 2023       |
|                     | <ul style="list-style-type: none"> <li>Developed curriculum and policy incorporating AI literacy and DEI principles</li> </ul>   |            |
|                     | <b>Networking Chair</b> , Yale Gruber Science Fellowship   | 2022, 2023 |
|                     | <ul style="list-style-type: none"> <li>Hosted networking talks, panels, and discussion to foster Gruber scientific community</li> </ul>  |            |
|                     | <b>Student Representative</b> , Graduate Student Assembly  | 2023       |
|                     | <ul style="list-style-type: none"> <li>Advocated for CB&amp;B graduate students to improve Yale healthcare literacy and policy</li> </ul>  |            |
|                     | <b>Reviewing Editor</b> , Yale Journal of Biology & Medicine   | 2023       |
|                     | <ul style="list-style-type: none"> <li>Managed manuscripts for <i>Big Data</i> issue, including inviting reviewers and making editorial decisions</li> </ul>   |            |
|                     | <b>Social Services &amp; Insurance Counseling</b> , HAVEN Free Clinic  | 2022, 2023 |
|                     | <ul style="list-style-type: none"> <li>Provided healthcare guidance and resources to uninsured New Haven residents</li> </ul>  |            |
|                     | <b>Cancer Biology Training Program</b> , Yale School of Medicine   | 2021-2023  |
|                     | <ul style="list-style-type: none"> <li>Completed certificate in cancer biology through additional translational coursework</li> <li>Shadowed Dr. Pamela Kunz and discussed clinical relevance of <i>in silico</i> cancer research</li> </ul> |            |