# Aarya Venkat

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# **EDUCATION**

#### UNIVERSITY OF GEORGIA

PHD BIOCHEMISTRY ARCS Foundation Scholar Future Faculty Fellow Dec 2023

#### **UC, SAN DIEGO**

MS CHEMISTRY June 2017

# **SKILLS**

#### **PROGRAMMING**

Python • CSS • Git • Bash • Docker

#### **BIOINFORMATICS**

#### Sequence Analysis

Metagenomic sourcing • Sequence alignment • Intron/Exon analysis • Clustering • ESM inference

#### Molecular Dynamics

MDAnalysis • Amber suite • GROMACS • Schrodinger • Forcefield parameterization

## Molecular Modeling

AlphaFold2 • Rosetta • Molecular Docking • VMD • PyMOL • Homology modeling • RFDiffusion

#### Deep Learning

Explainable AI • Protein language models • Dataset preparation • App/Tool development • Predictive modeling

#### **RESEARCH**

Writing • Editing • Analytical skills • Data Visualization • Figure generation • Illustrator • Multivariate Statistical Analysis • Making presentations

# **MISCELLANEOUS**

Leadership • Public speaking • Documentation • Systems administration

# **AWARDS**

2023 | Eriksson Lecture Scholarship

2023 | ASBMB Conference Travel Award

2022 | Glycobiology Poster Award

2022 | Glycobiology Travel Award

2021 | Glycobiology Poster Award

2021 | Glycobiology Travel Award

2021 | UGA Travel Award

2021 | ARCS Foundation Scholar

2021 | Three Minute Thesis Award

# RECENT EMPLOYMENT HISTORY

## GINKGO BIOWORKS, INC | PROTEIN ENGINEER III

Dec 2023 - current | Boston, MA

- Combined molecular docking/modeling, inference with evolutionary models like ESM2 and evcouplings, and metagenomic sourcing approaches to research and design or source a wide variety of proteins requested in high-budget contracts.
- Developed custom deep learning models fine-tuned from ESM2 and other models to build custom enzyme models.
- Developed pipelines for tools like RFDiffusion-AA, proteinMPNN, etc to be used for large-scale design of proteins.
- Subject matter expert and tech liaison for multiple projects involving Al, protein variant design, or general computational biology expertise.

# **UNIVERSITY OF GEORGIA** | GRADUATE RESEARCH ASSISTANT July 2018 - Dec 2023 | Athens, GA

- Performed bioinformatics and evolutionary analyses, including molecular dynamics, phylogeny, modeling, quantum chemistry, and sequence analyses.
- Developed deep learning models and tools to classify enzyme function.
- Maintain data storage, equipment, purchase orders, protocol documentation.
- Authored over 17 manuscripts for publication, with key involvement in writing, editing, data generation, analysis, visualization, and funding.

# UC, SAN DIEGO | GRADUATE RESEARCH ASSISTANT

Oct 2015 - May 2017 | La Jolla, CA

• Cytoscape intern at National Resource for Network Biology (2016). Developed PathInsight, an app to model the effects of small molecules.

# SELECTED PUBLICATIONS

Papers (Total: 21) (citations: 200) (H-index: 8)

<u>Venkat, A.</u> (2024). GapClean, a tool for cleaning and improving comprehension of protein sequence alignments. In Prep.

<u>Venkat, A.</u>, et al. (2024). Glydentify, a deep learning tool for classification of glycosyltransferase function. In Prep.

 $\underline{\text{Venkat}, A.} \text{ et al. (2023)}. \text{ Mechanistic and evolutionary insights into isoform-specific 'supercharging' in DCLK family kinases. eLife.}$ 

Aceil, J. $\dagger$ , <u>Venkat, A.</u> $\dagger$ , et al. (2023). Prevalence and homology of the pneumococcal serine-rich repeat protein at the global scale. Microbiology Spectrum.

Meng, Y. et al. (2023). Phosphorylation-dependent pseudokinase domain dimerization drives full-length MLKL oligomerization. Nature Communications

<u>Venkat, A.</u>, et al., (2022). Modularity of the hydrophobic core and evolution of functional diversity in fold A glycosyltransferases. Journal of Biological Chemistry.

Taujale, R., <u>Venkat, A.</u>, et al. (2020) Deep evolutionary analysis reveals the design principles of fold A glycosyltransferases. eLife.

# INVITED TALKS

#### 2023

Evolution of fold-A glycosyltransferases and GTXplorer, a new tool for comparative glycomics. ASBMB Conference. Seattle Conference Center

Evolution of functional diversity of fold A glycosyltransferases. Eriksson Lecture. Complex Carbohydrate Research Center

#### 2022

AlphaFold2: protein structure-prediction in the modern era. BCMB3600. University of Georgia

Modularity of the hydrophobic core and evolution of functional diversity in fold A glycosyltransferases. Southeast Enzyme Conference

#### 2017

Principles of chemical equilibria, UC, San Diego

# WORKSHOPS AND SYMPOSIA

#### 2023

Al and Deep Learning, Applications in Bioinformatics

#### 2022

Protein Engineering Symposium - Chair

#### 2021

Unfolded Protein Response Symposium - Chair

#### 2020

Writing a Diversity Statement - GradTeach Workshop - Workshop leader, 2020

Spring Teaching Symposium - Workshop leader

Spring Teaching Symposium - Organizer

#### 2019

Cancer Immunotherapies Symposium - Organizer

# **SERVICE**

#### NATIONAL HISTORY DAY JUDGE

2020-2022 | Athens, GA

Judged Performances and Documentaries for middle school and high school students. Provided critiques to aid them in state and national competitions.

#### SCIENCE OLYMPIAD LAB MANAGER

2017-2020 | Athens, GA

Aid in setting up Chemistry and Forensic labs for high school Science Olympiad. Ensure lab safety protocols are followed as students compete.

# PRESENTATIONS/POSTERS

#### 2023

Glydentify, a deep learning tool for classification of glycosyltransferase function. Society for Glycobiology. Big Island. HI

#### 2022

Modular evolution of fold-A glycosyltransferases and new tools to analyze them. Society for Glycobiology. Fernandina Beach, FL

Modularity of the hydrophobic core and evolution of functional diversity in fold A glycosyltransferases. Glycoscience Training Program. Athens, GA

#### 2021

Modularity of the hydrophobic core and evolution of functional diversity in fold A glycosyltransferases. Society for Glycobiology.

#### 2020

Playground Learning: Team Learning and Gamification. USG Teaching & Learning Conference (Cancelled: COVID-19)

Unfolded Protein Response. Biochemistry Symposium Chair. University of Georgia.

#### 2019

Deep Evolutionary Analysis Reveals the Design Principles of Fold A Glycosyltransferases. Glycobiology Conference Poster. PI: Natarajan Kannan, UGA

Cancer Immunotherapies. Biochemistry Symposium Host. University of Georgia.

Teaching and Laboratory Assistant Orientation. Delivered Lecture on Efficient Grading Practices. University of Georgia.

#### 2018

Teaching and Laboratory Assistant Orientation. "Teaching Tips" Q&A Panelist. University of Georgia.

#### 2017

PathInsight: A Novel Tool for Modeling Biomolecular Pathways. Thesis Defense. UCSD.

Guest Lecture on Chemical Equilibrium. Chemistry 6B, UCSD.

#### 2016

Does Competition Enhance Learning Over a Relaxed Guided Lesson? Teaching Methods Poster Presentation. Natural Sciences Building, UCSD.