

NIKIL KRISHNA

Undergraduate Student SASTRA Deemed University



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ABOUT ME

My academic and research interests lie at the intersection of biotechnology and computational biology, where I utilize digital tools to visualize and explore microscopic biological systems.

My experience spans molecular visualization, design, and simulation. I am presently deepening my understanding of artificial intelligence and machine learning applications in in silico drug design and protein binder design, with a particular focus on receptor pocket structure-based modeling.

Beyond academia, I enjoy listening to music, reading, and engaging in various forms of craftwork.

EDUCATION

2026

SASTRA Deemed University, Thanjavur **B.tech Biotechnology**

- 8.57 CGPA
- Honours
- Research Credits

2022

ALPHA Wisdom Vidyashram Tiruchirapalli

Higher Secondary Certificate

- Physics
- Chemistry
- Biology
- English
- Mathematics

2020

ALPHA Wisdom Vidyashram Tiruchirapalli

Secondary School Certificate

- English
- Hindi
- Science
- Mathematics
- Social Science

EXPERIENCE

2025

IIT Dharwad

Summer Internship

- Development of Early Stage Drug Discovery Pipeline Using Machine Learning Models
 - o Conda for virtual environemnt, dependency management
 - Python for workflow
- · Comparision of three SOTA machine learning architectures for protein binder
 - Bindcraft (Alphafold2 backpropagation)
 - BoltzDesign1 (Boltz1 backpropagation)
 - Rfdiffusion (Denoising diffusion)

2024

SASTRA Deemed University

Undergraduate Student Researher

- Molecular Motors Laboratory
- Drug Design and Protein Engineering

ML Protein Structure Prediction

- Alphafold
- ESMfold
- Chai-1
- Boltz
- Diffusion Based Models

Molecular Modelling

- ChimeraX
- Pymol
- VMD
- MOE
- Schrodinger Maestro

Drug Design

- · Virtual Screening
- Fragment Based Drug Design
- Genetic algorithm/Combinatorial Chemistry
- Lead Optimization
- ADMET analysis

Python

- Molecular Visualization
- Data Analysis
- Machine Learning
- Bioinformatics Pipelines
- Dependency management
- Al Automation

Protein Binder Design

- RFdiffusion
- ProteinMPNN
- Bindcraft
- ColabDesign
- Diffusion & Hallucination
 ML Models

Molecular Docking and

Dynamics

- Autodock
- Diffdock
- Gromacs
- NAMD
- OpenMM

Bioinformatics

- Linux
- Cryo-EM Modelling using ML
- BLAST, MSA
- Visual Studio Code
- Git version management
- Scientific Computing

Process Simulation

- Aspen Plus
- SuperPro Designer
- DWSIM

CERTIFICATIONS

2025

SASTRA Deemed

University

Aspen Plus Simulation Software - A Basic Course For Beginners

NPTEL • 90%

2024 Schrodinger Drug Discovery Hackathon Winner

To design a insilico de-novo structure-based drug molecule aimed at executing the dual inhibition of critical protein kinases A and B in Mycobacterium tuberculosis by genetic algorithm and combinatorial chemistry.

2024 Drug Delivery: Principles and Engineering

NPTEL • 83%

PROJECTS

2025 Founder - Biostruct

A web platform which provides computational biology tools in an

interactive online interface https://biostruct.netlify.app/

2024 Research Credits

SASTRA Deemed University

Fragment Based Drug Design of Anthelmintic Using Genetic Algorithm

LANGUAGES

English - Professional working proficiency
 Hindi - Limited working proficiency
 Tamil - Bilingual proficiency
 Kannada - Bilingual proficiency
 Telugu - Elementary proficiency

REFERENCES

Dr.Venkatasubramanian
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Superviser Associate Professor SASTRA Deemed University Mentor
Senior Assistant
Professor
SASTRA Deemed
University

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