Aditya Sarkar | C

■ asnov2k@gmail.com 🚱 aditya-sarkar441.github.io 🞧 aditya-sarkar441

Undergraduate Student at IIT Mandi

ACADEMIC HISTORY

University of California Los Angeles Junior Research Scientist

LOS ANGELES, CA June 2022 – August 2022

• Hosted by Sriram Sankararaman [Report]

• Sponsored by UCLA CS Department.

Indian Institute of Technology (IIT) Mandi Bachelor of Technology (Honors) in Electrical Engineering

KAMAND, HP July 2019 - May 2023

- Advised by Dileep A.D. [Thesis]
- CGPA of 9.1/10 (Inst. rank: 2, Dept. rank: 1)
- Supported by NTS Scholarship 2017-2023

AWARDS AND ACHIEVEMENTS

- Ranked first in the department (out of 60 students) with a CPI of 9.1/10.0 after 6th semester.
- Received IIT academic award for being department topper in Fall 2019 and Spring 2020.
- Awarded department change due to exceptional academic performance in the first year (9.8 CPI).
- Selected for attending the Winter School for Cognitive Modelling 2020 on Cognitive Modelling, organised by University of Waterloo, Canada and University of Groningen, Netherlands.
- Ranked in top 1% (amongst 2.2 Million students) in the entrance exam of Indian Institute of Technology (IIT-JEE) 2019 (99.3 percentile)
- Awarded National Talent Search Scholarship in 2017 by NCERT.

Publications and Preprints

2. Sarkar A., Joseph A.T., Muralikrishna H., A.D. Dileep LA-PERC-NET: Gated Perceptron Network for reducing the complexity of spoken language identification systems., To be submitted in Pattern Recognition Letters.

Journal.....

1. Sarwal V, Niehus S, Ayyala R, Kim M*, Sarkar A*, Chang S, Lu A, Rajkumar N, Darfci-Maher N, Littman R. Chhugani K. Sovlev A. Comarova Z. Wesel E. Castellanos J. Chikka R. Distler MG, Eskin E. Flint J. Mangul S., A comprehensive benchmarking of WGS-based deletion structural variant callers., in Briefings in Bioinformatics (IF=12). doi: 10.1093/bib/bbac221.

- 2. Sarkar A., Joseph A.T., Muralikrishna H., A.D. Dileep MHA-NET: Multi-Head Attention Network for improving the performance of spoken language identification systems in Real-World conditions., Submitted to IEEE Machine Learning for Signal Processing.
- 1. Sarkar A., Bhaysar A., Virtual Screening of Pharmaceutical Compounds with hERG inhibitory activity, in International Joint Conference on Biomedical Engineering Systems and Technologies, doi: 10.5220/0010267701520159.

TEACHING EXPERIENCE

Served as undergraduate teaching assistant for certain courses, conducting weekly tutorial sessions, special doubt sessions, and grading answer sheets for the following courses:

• CS671: Deep Learning and Applications (Prof. Dileep A.D.)

Summer 2023

• IC272: Introduction to Machine Learning (Prof. Aditya Nigam)

Fall 2022

• IC252: Probability and Statistics (Prof. Satyajit Thakur)

Summer 2022

• CXPT599: Biomedical Data Science (Prof. Serghei Mangul)

Summer 2022

Selected Courses

Advanced Courses: Tensors*, Convex Optimization*, Statistical Learning, Deep Learning

Core EE courses: Digital Communication*, Control Systems*, Digital Systems Design*, Digital Image Processing. Computer Science: Introduction to Machine Learning, Data Structures and Algorithms, Data Visualization, Data

Science, Computer Networks, Computer Organization

Mathematics: Probability and Statistics, Calculus, Linear algebra, Matrix theory*

OLDER RESEARCH PROJECTS

Deep Cross Modality Hashing using Cycle Consistent Transformer Guide: Aditya Nigam

Sept '22 - Dec '22

IIT Mandi

- Developed a cyclic end-to-end transformers that can form text representations from image, and image representations from text so as to reduce the modality gap. Loss function used was L2-Norm.
- Used ArcFace loss function to perform unimodal hashing over image representations of both the images and text. Currently exploring the developed model over various datasets such as NUS-Wide, Mir-Flickr25K and MS-COCO, trying to get better results over a variety of metrics. [Slides]

Cell type decomposition using Expectation Maximization

Jun '22 - Oct '22

Guide: Serghei Mangul

University of Southern California

- Surveyed and implemented existing models on deconvoluting cell types proportions from gene expression data. Used GEDIT's feature selection process based on information content to select the most probable set.
- Developed an Expectation Maximization based algorithm for estimating the cell type proportion from bulk gene expression data and reference data. [Report]

TECHNICAL SKILLS

Languages: C++, Python, R, VHDL, MATLAB, Bash, HTML/CSS, Javascript

Deep Learning: Caffe, TensorFlow, Numpy, Pandas, OpenCV, Matplotlib, Keras, Pytorch

Software: Keil μ Vision, LATEX, Git, Scilab, SolidWorks, Arduino, Raspberry Pi, Modelsim, UCLA Hoffman2 Cluster