Souvik Ghosh

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About me

I am Souvik Ghosh, a motivated and detail-oriented software developer and researcher, currently pursuing an M.Sc. in Computer Science and Engineering at BUET. I am passionate about software development, machine learning, and problem-solving. I am eager to leverage my skills in research and development while contributing effectively to academia and industry.

Education _____

MSc. Bangladesh University of Engineering and Technology, Computer Science

Feb 2025 - Present

- · Current Graduate Student
- Research areas: Bioinformatics, Machine Learning, Deep Learning

BSc. Bangladesh University of Engineering and Technology, Computer Science

Feb 2020 - Feb 2025

- · CGPA: 3.85/4.0
- **Relevant Coursework:** Compilers, Operating Systems, Computer Networks, Database, DSA, Software Engineering, Information System Design

Experience _____

BRAC University, Lecturer, CSE Department

Jun 2025 - Present

- Teaching undergraduate courses in CSE and mentoring student projects.
- Continuing research in bioinformatics and deep learning applications.

Apurba Technologies, Full Stack Developer

Mar 2025 - May 2025

- Developed full-stack applications with ReactJS, Node.js, and PostgreSQL.
- Built scalable REST APIs and optimized CI/CD pipelines with Docker and GitHub Actions.

Publications _____

ResLysEmbed: A ResNet-Based Framework for Succinylated Lysine Residue Prediction Using Sequence and Language Model Embeddings

2025

Souvik Ghosh, Md Muhaiminul Islam Nafi, Mohammad Saifur Rahman Bioinformatics Advances, vbaf198 (2025) ☑

Research Experience _____

Generic Post-Translational Modification (PTM) Site Detection

Ongoing

Supervisor: Dr. Swakkhar Shatabda, Professor, BRAC University

- Developing a general framework for PTM site prediction using protein language model embeddings combined with structural data.
- Modeling proteins as graphs where each amino acid is a node initialized with PLMbased embeddings.
- Exploring Graph Neural Networks (GNNs) and related architectures for effective sequence–structure integration.

 Aiming for a unified approach that can be adapted to multiple PTM types instead of task-specific models.

Alignment-Free Phylogenetic Tree Construction using Syncmers and TF-IDF

Ongoing

Supervisor: Dr. Atif Hasan Rahman, Professor, BUET CSE

- Investigating a novel alignment-free method for phylogenetic inference by replacing traditional *k*-mers with **syncmers**.
- Introducing **TF-IDF scoring**, inspired by natural language processing, instead of raw frequency or binary presence/absence for sequence representation.
- Constructing syncmer TF-IDF matrices to capture discriminative sequence features.
- Exploring machine learning approaches to train on these representations for accurate phylogenetic tree reconstruction.

Undergraduate Thesis, CSE, BUET

Published 2025

Title: ResLysEmbed: A ResNet-Based Framework for Succinylated Lysine Residue Prediction Using Sequence and Language Model Embeddings

Supervisor: Dr. Mohammad Saifur Rahman

- Published in Bioinformatics Advances (2025).
- Achieved state-of-the-art performance in succinylation site prediction, surpassing existing benchmarks.

Deepfake Detection for IEEE SP Cup 2024

Submitted

Supervisor: Dr. Mohammad Saifur Rahman

- Developed hybrid deep learning models for generalizable deepfake detection.
- Explored generative approaches, including diffusion models, GANs, and VAEs for dataset augmentation.
- Evaluated advanced backbone architectures, such as MaxViT, ConvNeXt, and EfficientNet, with specialized loss functions like contrastive loss and focal loss for improved accuracy.

Projects _

Bits Unplugged | Interactive CS Learning Platform

- Co-developed an innovative platform focused on enhancing problem-solving skills without coding, emphasizing strategy over syntax.
- Implemented features such as drag-and-drop interactive problem solving, realtime contests, personalized recommendations, and analytics dashboards.
- Technologies: Docker, GitHub Actions, Tailwind CSS, Sequelize, PostgreSQL, Supabase, ReactJS, NodeJS, ExpressJS, Render.com

Multimodal Breast Cancer Prognosis Prediction | Machine Learning Project

Github 🗹

Website 🗹

Github ☑ YouTube ☑

- Designed a multimodal framework using mRNA expression, copy number alteration, and clinical data from the TCGA-BRCA dataset.
- Transitioned from MLP-based models to self-attention and cross-attention mechanisms for improved performance.
- Focused on contrastive learning techniques to handle feature embeddings without classification during training.

Awards & Achievements _

Research Grant from RISE BUET

Received a research grant from BUET Research and Innovation Centre for Science and Engineering for an ongoing research project.

Deepfake Detection for IEEE SP Cup 2024

Selected as a finalist in the top 3 teams; the final competition will occur in April 2025.