Md Muhaiminul Islam Nafi

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OBJECTIVE

I am Md. Muhaiminul Islam Nafi. I am currently an undergrad student of Computer Science at BUET. I want to build a successful career using my technological knowledge and relevant skills. I want to contribute to the betterment of the place I am allowed to work for. I also want to hone my other skills like time management, fast adaptability, and creativity.

MOTIJHEEL GOVT. BOYS' HIGH SCHOOL 2015-2017

SSC — GPA: 5 00/5 00

NOTRE DAME COLLEGE DHAKA 2017-2019

HSC - GPA: 5.00/5.00

BANGLADESH UNIVERSITY OF ENGINEERING AND TECHNOLOGY 2020-present

B.Sc. in Computer Science and Engineering — CGPA: 3.96/4.00 (up to 7 terms)

SKILLS

PROGRAMMING LANGUAGES Python | Javascript | Typescript | C | c++ | Assembly (x86, MIPS) | Bash | Java | LaTeX WEB DEVELOPMENT HTML | CSS | Express | React | Svelte | Django | Next.js | Figma | Docker | Spring Boot

Matplotlib | NumPy | Pandas | Scikit-learn | PyTorch MACHINE LEARNING

TOOLS AND TECHNOLOGIES Design Patterns | Git | Microservices architecture | Swagger API | Postman | MISP | Markdown

> DATABASES MySQL | PostgreSQL | Oracle | Prisma ORM

WordPress CONTENT MANAGEMENT GAME DEVELOPMENT Pygame | Unity

PUBLICATIONS

Prediction of protein-carbohydrate binding sites from protein primary sequence (Under review)

In this study, we propose StackCBEmbed, an ensemble machine learning model to effectively classify protein-carbohydrate binding interactions at the residue level. StackCBEmbed combines traditional sequence-based features along with features derived from a pretrained transformer-based protein language model. Link to the initial version of the manuscript: bioRxiv

DeepBCTPred: Deep Learning-Based Prediction of Bladder Cancer Tissues from Endoscopic Images (Under review)

In this study, we designed a pipeline to generate new images and a novel genetic algorithm to effectively select images from them. Additionally, we combined handcrafted features with learned features.

Predicting Protein-Carbohydrate Binding Sites: A Deep Learning Approach Integrating Protein Language Model Embeddings and **Structural Features** (Manuscript in preparation)

In this study, we created a novel deep learning architecture that integrates various dataset imbalance approaches with structural features.

AWARDS AND HONORS

BUET RISE GRANT

RISE Student Research Grant [No. S2024-01-004] (GRANT RECEIVE)

HONORABLE MENTIONS MicroProcessor and MicroController project

PROJECTS

ONCAMPUS

GITHUB LINK: BACKEND FRONTEND

Frontend: Framework: Next.js, Styling: Tailwind CSS, Component Library: Material Tailwind, Text Box: jodit-react, PDF Viewer: react-pdf, 360° Virtual Tour: react-photo-sphereviewer, Language: TypeScript

Backend: Framework: Node.js, Express, Architecture: Microservice Architecture, ORM: Prisma ORM, Security: Helmet, JWT, Authentication: Keycloak, NextAuth.js. Database: PostgreSQL hosted on Supabase, File Storage: Edgestore, Google Calendar Integration: Google Cloud API.

Deployment: MS Azure Virtual Machines, Supabase, Docker (for Keycloak), SSL Certificate from Namecheap, API Documentation with Postman.

MOOMARKET

GITHUB LINK: BACKEND & FRONTEND

Tech Stack: JavaScript (Node.js, Vanilla JS), Express, HTML, EJS, PostqreSQL, Git, GitHub, npm, Render.

GITHUB LINK: BACKEND & FRONTEND

Tech Stack: JavaScript (Node.js, Vanilla JS), Express, HTML, EJS, Oracle DB, Git, GitHub, npm.

CERTIFICATES

PERFECT ATTENDANCE CERTIFICATE Issued May 2020

NOTRE DAME COLLEGE DHAKA CERTIFICATE IN NATIONAL SKILL STANDARD BASIC COURSE EXAMINATION, 2015 Issued Sep 2015

BANGLADESH TECHNICAL EDUCATION BOARD

INTERESTS

- · Watching movies
- · Reading novels
- Listening to music