

# Kh Mujahidul Islam

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## Education

### Bachelor of Science (B.Sc.)

Feb 2025

Department of Computer Science and Engineering,  
Faculty of Engineering and Technology, Islamic University, Bangladesh.

CGPA: 3.27 / 4

Total Credit hours: 160

## Research Experience

### Research Assistant

Jan 2023 – Present

CABAIR (Center for Advanced Bioinformatics and AI Research), Islamic University, Bangladesh.

[Website](#)

### Single cell RNA sequencing project:

#### 1. Cell-specific biomarker identification in intellectual disability (Published in Scientific Reports):

- Conducted single-cell RNA sequencing (scRNA-seq) analysis using Seurat (PCA, UMAP for dimensionality reduction and clustering).
- Built protein–protein interaction (PPI) networks to explore gene interactions and performed ROC analysis for biomarker diagnostic evaluation.

#### 2. NK cell-based Biomarker Identification of Prostate Cancer (Manuscript in preparation):

- Conducted bulk RNA-seq and single-cell RNA-seq analyses to identify NK cell-specific biomarkers for prostate cancer, including cell clustering, differential expression, hub gene identification, pathway enrichment, and survival analysis.
- Performed virtual screening and molecular docking of hub genes.

### Weighted Gene Co-expression Network Analysis (WGCNA) Project:

#### Identification of potential biomarkers of Kidney Renal Clear Cell Carcinoma (KIRC) using WGCNA and development of machine learning-based staging models (Manuscript in preparation) :

- Applied WGCNA on TCGA datasets to identify co-expression modules and develop machine learning-based staging models.
- Used Lasso regression for feature selection and validated model performance with classification metrics.
- Identified potential biomarkers for Kidney Renal Clear Cell Carcinoma (KIRC), integrating bioinformatics and ML approaches for translational research.

Supervised by: **Dr. Md Habibur Rahman**

(email: [habib@iu.ac.bd](mailto:habib@iu.ac.bd))

## Professional Experience

### Intern Engineer - AI Agent Development

Jul 2025 – Present

Next Solution Lab, Mirpur DOHS, Dhaka.

- Contributed to the development of prototype AI agents and conversational pipelines with LangChain, LangGraph, Llama, and Ollama, enabling automated data retrieval and improved user engagement.
- Implemented model training and fine-tuning workflows; prepared datasets and evaluation scripts to measure model behaviour.
- Packaged prototypes as lightweight services (Flask/Streamlit) for internal demonstration and stakeholder review.
- Documented integration points and created demo notebooks to accelerate handover to engineering teams

## Notable Publications

**KM Islam**, MH Rahman, MA Hossain, M Khan, MM Rahman, PS Apu, MI Hasan, SA Ansari, & M Humayoo. **May 2025**

*Identification of cell-specific biomarkers for intellectual disability via single-cell RNA sequencing and transcriptomic bioinformatics approaches.* Scientific Reports. [DOI:10.1038/s41598-025-85162-4](https://doi.org/10.1038/s41598-025-85162-4) (IF:3.9)

MM Rahman, MH Rahman, MA Hossain, **KM Islam**, PS Apu, M Khan, MG Kibria, SA Ansari, & M Humayoo. **May 2025**

*Uncovering dendritic cell-specific biomarkers for diagnosis and prognosis of cardiomyopathy using single-cell RNA sequencing and comprehensive bioinformatics analysis.* Scientific Reports.

[DOI:10.1038/s41598-024-78011-3](https://doi.org/10.1038/s41598-024-78011-3) (IF:3.9)

MI Hasan, **KM Islam**, MN Hasan, MM Rahman, MH Rahman, S Moon, MN Park, HN Kang, & B Kim. **Mar 2025**

*Exploring the therapeutic potential of Evodia rutaecarpa in early-onset pancreatic cancer: A network pharmacology and molecular docking approach.* Current Pharmaceutical Design.

DOI: [10.2174/0113816128356225250305081937](https://doi.org/10.2174/0113816128356225250305081937) (IF 2.8)

M Khan, MA Hossain, **KM Islam**, MM Rahman, PS Apu, IA Ansari, SA Ansari, & MH Rahman. **Feb 2025**

*Integrative analysis of single-cell and bulk RNA sequencing data reveals T-cell specific biomarkers for diagnosis and assessment of celiac disease: A comprehensive bioinformatics approach.* Current Bioinformatics.

DOI: [10.2174/0115748936353313250123071744](https://doi.org/10.2174/0115748936353313250123071744) (IF 2.9)

## Technical Skills

<b>Programming and scripting</b>	Python , R , C++ , JavaScript.
<b>ML / DL skills</b>	Scikit-learn, Pandas, NumPy, Matplotlib, Seaborn, TensorFlow, Keras, PyTorch.
<b>Dry Lab Skills</b>	Microarray data analysis, scRNA sequencing data analysis, Weighted Gene Co-expression Network Analysis (WGCNA), Virtual Screening, Molecular Docking.
<b>Tools and Software</b>	Discovery Studio, PyMol, PyRx, AutoDock Vina, Cytoscape, DAVID, Network Analyst, BiocManager, Biopython, Mendeley, EndNote, LaTeX
<b>Databases</b>	PostgreSQL, NCBI, Ensembl, UniProt, PDB, GEO, TCGA.

## STANDARDIZED LANGUAGE TEST

*International English Language Testing System (IELTS) Academic*

TRF number: 25BD505875ISLK001A

Module	Listening	Reading	Speaking	Writing	Overall
Band Score	8.0	7.0	6.5	6.0	7.0

## Additional Informations

### Organizations :

<b>Executive Member</b> , IEEE Computer Society, IU SB, Islamic University, Bangladesh	<b>2022-2023</b>
<b>Additional treasurer</b> , Islamic University IT Society (IUIITS), Islamic University, Bangladesh	<b>2021-2022</b>
<b>Additional Secretary</b> , Tangail Zilla Student Welfare Association, Islamic University, Bangladesh	<b>2023-2024</b>
<b>Executive Member</b> , Cancer Awareness Program (CAP), Islamic University, Bangladesh	<b>2019-2020</b>
<b>Founding Member</b> , Help For Distressed People, MSSHSS, Tangail, Bangladesh.	<b>2016-present</b>

## References

### 1. Dr. Md Habibur Rahman

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### 2. Md. Ibrahim Abdullah

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### 3. Dr. Md. Robiul Hoque

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