



SAEED AHMED

📍 **Home** : Village and post office nawagai tehsil mandnr district buner, 19300, Buner, Pakistan

📍 **Work** : Department of Computer Science, University of Swabi, Pakistan, 23570, Swabi, Pakistan

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🌐 **Website**: <https://saeed-website.vercel.app/> 🗨 **WeChat**: saeedbuner

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📘 **Facebook**: <https://www.facebook.com/> 🐦 **X**: <https://twitter.com/home>

🌐 **LinkedIn**: <https://www.linkedin.com/mynetwork/>

👤 **sciprofiles**: <https://sciprofiles.com/profile/390359>

Gender: Male **Date of birth**: 02/02/1987 **Nationality**: Pakistani

WORK EXPERIENCE

Center for Research Innovation and Biomedical Informatics, Faculty of Medical Technology, MU

City: Bangkok | **Country**: Thailand

[01/07/2024 – Current] **Postdoctoral Research Fellow**

- Algorithm Development.
- Data Analysis
- Model Validation and Evaluation
- Collaboration
- Publication and Dissemination

Department of Computer Science, University of Swabi, Khyber Pakhtunkhwa, Pakistan

City: Swabi | **Country**: Pakistan

[30/09/2024 – Current] **Assistant Professor in Computer Science**

- Teaching
- Research

Department of Experimental Medical Science Lund University BMC B13 SE-22184 Lund Sweden

City: Lund | **Country**: Sweden

[10/09/2023 – 07/11/2024] **Postdoctoral Researcher**

- Research
- Algorithm Development.
- Data Analysis
- Model Validation and Evaluation
- Collaboration
- Publication and Dissemination

University Management and Technology Lahore

Address: C-II Block C 2 Phase 1 Johar Town, Lahore, Punjab 54770, Pakistan, University of Management and Technology Lahore, C-II Block C 2 Phase 1 Johar Tow, 54770, Lahore, Pakistan

[02/09/2021 – 31/08/2023] **Assistant Professor in Computer Science**

- Teaching
- Research
- Advising Students
- Supervising Student Projects, MS Thesis.

Center of data mining and biomedical informatics, Mahidol University, Bangkok Thailand,

City: Bangkok | **Country:** Thailand

[01/06/2021 – 28/02/2022] **Postdoctoral Research Fellowship**

- Research
- Algorithm Development.
- Data Analysis
- Model Validation and Evaluation
- Collaboration
- Publication and Dissemination

Higher Education Department KPK, Govt: Degree Collage Daggar Buner, KPK

[05/10/2020 – 31/08/2021] **Lecturer in Computer Science**

EDUCATION AND TRAINING

[08/10/2016 – 25/03/2021] **PhD in Computer Science and Technology (Bioinformatics)**

Nanjing University of science and technology Nanjing China <https://english.njust.edu.cn/>

Address: 200 xiaolingwei, Xuanwu district, jiangsu province, 210094, Nanjing, China |

[03/03/2013 – 28/09/2016] **MS in Computer Science**

Abdul Wali Khan University Mardan, KPK, Pakistan <https://awkum.edu.pk/>

Address: Bacha Khan Monument, Nowshera Mardan Rd, Muslimabad, Mardan, Khyber Pakhtunkhwa, 23200, Mardan, Pakistan |

[08/10/2007 – 28/12/2011] **BS in Telecommunication**

Institute of Engineering and Technology, University of Science and Technology Bannu KPK, Pakistan <https://www.ustb.edu.pk>

Address: Bannu, Khyber Pakhtunkhwa, 28210, Bannu, Pakistan |

[01/08/2004 – 13/07/2006] **Fsc Pre-Engineering**

Board of Intermediate and Secondary Education Swat, KPK Pakistan <https://bisess.result.pk>

Address: Kukrai, Saidu Sharif, Swat, Khyber Pakhtunkhwa , 19200, Swat, Pakistan |

[04/10/2000 – 18/07/2004] **Matriculation**

Board of Intermediate and Secondary Education Swat, KPK Pakistan <https://www.bisess.edu.pk>

Address: Kukrai, Saidu Sharif, Swat, Khyber Pakhtunkhwa , 19200, Swat, Pakistan |

THESIS/DISSERTATIONS

Research on Prediction of Protein Phosphorylation and Cancer Subtypes via Intelligent Computation.

Ph.D. Dissertation:

Submitted to Nanjing University of Science and Technology, Nanjing, China.

Identification of Heat Shock Protein Families and J-Protein Types using Dipeptide Composition and Support Vector Machine.

MS Thesis: Submitted to Abdul Wali Khan University, Mardan, Khyber Pakhtunkhwa, Pakistan.

CAREER OBJECTIVE AND RESEARCH INTEREST

Career Objectives

To join a dynamic and fast-growing organization in a challenging position where my educational qualifications, research experience, exposure, and strong work ethic can be fully utilized. I am a self-starter—self-driven, self-disciplined, enthusiastic, loyal, and fully committed to my employer.

Research Interest

To advance the development of cutting-edge computational tools and predictive models using machine learning and deep learning techniques for the analysis of protein, peptide, and genomic sequences. My goal is to contribute to the fields of **bioinformatics** and **bio medical informatics** by designing interpretable, accurate, and efficient algorithms for biomolecular function prediction, therapeutic peptide discovery, and disease biomarker identification. I aim to integrate multi-omics data and AI-driven approaches to address complex biological problems and support precision medicine initiatives.

LANGUAGE SKILLS

Mother tongue(s): Pashto

Other language(s):

Urdu

LISTENING C2 READING C2 WRITING C2

SPOKEN PRODUCTION C2 SPOKEN INTERACTION C2

English

LISTENING C2 READING C2 WRITING C2

SPOKEN PRODUCTION C2 SPOKEN INTERACTION C2

Chinese

LISTENING A1 READING A1 WRITING A1

SPOKEN PRODUCTION A1 SPOKEN INTERACTION A1

Levels: A1 and A2: Basic user; B1 and B2: Independent user; C1 and C2: Proficient user

SKILLS

Bioinformatics and Computational Biology | Artificial Intelligence | Deep Learning | pattern recognition | Computational biology | Python/PyTorch | Knowledge of C, Python, Java, Matlab | Python libraries (Numpy Pandas Matplotlib Tensorflow networkx) | SLURM HPC | Python (matplotlib numpy scipy pandas scikit-learn tensorflow) | C, C#, C++ | HPC clusters (Slurm) | TensorFlow | Multi-modal biological and biomedical data integration | Multi-modal | Python | Machine Learning | Scikit-Learn | Keras | Git | artificial intelligence

PUBLICATIONS

[2025]

[Accurate identification of broadly neutralizing antibodies against dengue virus based on deep stacking strategy with multi-perspective features](#)

Reference: Ahmed, S., Schaduengrat, N., Pipattanaboon, C. et al. Accurate identification of broadly neutralizing antibodies against dengue virus based on deep stacking strategy

with multi-perspective features. Sci Rep (2025).

Authors: S Ahmed, N Schaduangrat, C Pipattanaboon & W Shoombuatong | **Journal Name:** Scientific Reports | **Publisher:** Nature Publishing Group UK

[2025] [**BLSAM-TIP: Improved and robust identification of tyrosinase inhibitory peptides by integrating bidirectional LSTM with self-attention mechanism**](#)

Reference: October 8, 2025

Authors: S Ahmed,N Schaduangrat,P Chumnnpuen,S. M. H Mahmud,K O M Goh,W Shoombuatong | **Volume, Issue and Pages:** 20,10,e0333614 | **Publisher:** Public Library of Science

[2025] [**DeepHDAC3i: Leveraging an interpretable deep learning-based framework for the accelerated discovery of HDAC3 inhibitors**](#)

Reference: IEEE/ACM Transactions on Computational Biology and Bioinformatics

Authors: Saeed Ahmed, Nalini Schaduangrat, Ittipat Meewan, Watshara Shoombuatong | **Journal Name:** IEEE/ACM Transactions on Computational Biology and Bioinformatics | **Volume, Issue and Pages:** p 1-13, pubMed ID 40880325 | **Publisher:** IEEE

[2025] [**PON-P3: Accurate Prediction of Pathogenicity of Amino Acid Substitutions**](#)

Authors: Muhammad Kabir, Saeed Ahmed, Haoyang Zhang, Ignacio Rodríguez-Rodríguez, Seyed Morteza Najibi, Mauno Vihinen | **Journal Name:** International Journal of Molecular Sciences | **Volume, Issue and Pages:** 26,5,2004 | **Publisher:** MDPI

[2025] [**Advancing the Accuracy of Anti-MRSA Peptide Prediction Through Integrating Multi-Source Protein Language Models**](#)

Authors: Watshara Shoombuatong, Nalini Schaduangrat, Nutta Homdee, Saeed Ahmed, Pramote Chumnnpuen | **Journal Name:** Scientific Reports | **Volume, Issue and Pages:** 15,1,4762 | **Publisher:** Nature Publishing Group UK

[2025] [**Advancing the Accuracy of Anti-MRSA Peptide Prediction Through Integrating Multi-Source Protein Language Models**](#)

Authors: Watshara Shoombuatong, Pakpoom Mookdarsanit, Lawankorn Mookdarsanit, Nalini Schaduangrat, Saeed Ahmed, Muhammad Kabir, Pramote Chumnnpuen | **Journal Name:** Interdisciplinary Sciences: Computational Life Sciences | **Volume, Issue and Pages:** 1-14 | **Publisher:** Springer Nature Link

[2025] [**A comprehensive review and evaluation of machine learning-based approaches for identifying tumor T cell antigens**](#)

Authors: Watshara Shoombuatong, Saeed Ahmed, SM Hasan Mahmud, Nalini Schaduangrat | **Journal Name:** Computational Biology and Chemistry | **Volume, Issue and Pages:** 118 | **Publisher:** 108440

[2025] [**GRU4ACE: Enhancing ACE inhibitory peptide prediction by integrating gated recurrent unit with multi-source feature embeddings**](#)

Reference: Ahmed, S., Schaduangrat, N., Chumnnpuen, P., & Shoombuatong, W. (2025). GRU4ACE: Enhancing ACE inhibitory peptide prediction by integrating gated recurrent unit with multi-source feature embeddings. Protein Science, 34(6), e70026.

Authors: Saeed, A., Schaduangrat, N., Chumnnpuen, P., & Shoombuatong, W. | **Volume, Issue and Pages:** 34,6 | **Publisher:** John Wiley & Sons, Inc.

[2024] [**A Deep Learning Model for Detecting Fake Medical Images to Mitigate Financial Insurance Fraud**](#)

Reference: M A Arshed, S Mumtaz, S tefan C Gherghina, N Urooj, S Ahmed, and C Dewi; Computation 2024, 12, 173.

- [2024] [**A novel deep learning identifier for promoters and their strength using heterogeneous features**](#)
- Reference:** A Amjad, S Ahmed, MKabir, M Arif, T Alam, 19 August 2024, Methods
- [2024] [**A Computational Predictor for Accurate Identification of Tumor Homing Peptides by Integrating Sequential and Deep BiLSTM Features**](#)
- Reference:** R Arif, S Kanwal, S Ahmed & MKabir, 11 May 2024, Vol 16, pages 503–518, Computational Life Sciences
- [2024] [**A Comparative Review and Analysis of Computational Predictors for Identification of Enhancer and their Strength**](#)
- Reference:** M Gill, M Kabir, S Ahmed, M A Subhani, and MHayat;Current Bioinformatics, xxxx, XX, x-x
- [2024] [**An explainable stacking-based approach for accelerating the prediction of anti-diabetic peptides**](#)
- Reference:** F Arshad, S Ahmed, A Amjad, and M Kabir; 25 April 2024, 115546; Analytical Biochemistry
- [2024] [**A novel stacking-based predictor for accurate prediction of antimicrobial peptides**](#)
- Reference:** S Kanwal, R Arif, S Ahmed and M Kabir; Journal of Biomolecular Structure and Dynamics [J], 1-12
- [2024] [**A 16 × 16 Patch-Based Deep Learning Model for the Early Prognosis of Monkeypox from Skin Color Images**](#)
- Reference:** M A Arshed,H A, Rehman, S Ahmed, C Dewi, and H J Christanto; Computation 2024, 12(2), 33
- [2024] [**Multiclass AI-Generated Deepfake Face Detection Using Patch-Wise Deep Learning Model**](#)
- Reference:** M. A. Arshed, S. Mumtaz,M. Ibrahim,C. Dewi ,M. Tanveer and S. Ahmed
- [2023] [**A Novel Predictor for the Analysis and Prediction of Enhancers and Their Strength via Multi-View Features and Deep Forest**](#)
- Reference:** M Gill,S Ahmed,M Kabir, and M Hayat; Information 14.12 (2023): 636.
- [2023] [**Chem2Side: A Deep Learning Model with Ensemble Augmentation \(Conventional+ Pix2Pix\) for COVID-19 Drug Side-Effects Prediction from Chemical Images**](#)
- Reference:** M A Arshed,M Ibrahim,S Mumtaz, M Tanveer and S Ahmed, Information 14.12 (2023): 663.
- [2023] [**Multi-class skin cancer classification using vision transformer networks and convolutional neural network-based pre-trained models**](#)
- Reference:** Arshed, M. A., Mumtaz, S., Ibrahim, M., Ahmed, S., Tahir, M., & Shafi, M..Information,14(7), 415.
- [2023] [**IoT Based Smart Baby Monitoring System with Emotion Recognition Using Machine Learning**](#)
- Hina Alam, Muhammad Burhan, Anusha Gillani, Ihtisham ul Haq, Muhammad Asad Arshed, Muhammad Shafi, and **Saeed Ahmad**

- [2022] [PredAoDP: Accurate Identification of Antioxidant Proteins by Fusing Different Descriptors Based on Evolutionary Information with Support vector machine. Accepted, \[J\] Chemometrics and Intelligent Laboratory Systems.](#)
S Ahmed, M. Kabir*, M. Arif, K. Khan, Y. D Khan;
- [2022] [AMY-FRL: a novel sequence-based approach for accurate prediction of amyloid proteins by using feature representation learning. \[J\] Scientific Report.](#)
P Charoenkwan, S Ahmed, C Nantasenamat, W Shoombuatong,*;
- [2022] [SCORPION is a stacking-based ensemble-learning framework for accurate prediction of phage Virion proteins. \[J\] Scientific Report](#)
S Ahmed, P Charoenkwan, J M.W. Quinn, M A Moni, Md Mehedi Hasan, Pietro Lio' and W Shoombuatong;
- [2022] [StackACPred: prediction of anticancer peptides by integrating optimized multiple feature descriptors with stacked ensemble approach. \[J\] Chemometrics and Intelligent Laboratory Systems](#)
M Arif, S Ahmed, M Kabir, and D-J Yu*
- [2021] [DeepCPPred: a deep learning framework for discrimination of cell-penetrating peptides and their uptake efficiencies. \[J\] IEEE/ACM Transactions on Computational Biology and Bioinformatics.](#)
M Arif, M Kabir*, S Ahmed, A Khan, G Feng, A Khelifi, DJ Yu;
- [2021] [DeepPPSite: A deep learning-based model for analysis and prediction of phosphorylation sites using efficient sequence information. \[J\] Analytical Biochemistry.](#)
S Ahmed, M Kabir, M Arif, Z U Khan and DJ Yu*;
- [2020] [TargetCPP: accurate prediction of cell-penetrating peptides from optimized multi-scale features using gradient boost decision tree.\[J\] Journal of Computer-Aided Molecular Design.](#)
M Arif, S Ahmad, F Ali, G Fang, M Li, DJ Yu;
- [2020] [HPhosPPred: Prediction of human phosphorylated proteins by extracting discriminative features from the evolutionary profile and physicochemical properties through LFDA. \[J\] Chemometrics and Intelligent Laboratory Systems](#)
S Ahmad*, M Kabir*, M Arif, Z Ali and ZNK Swati;
- [2020] [Pred-BVP-Unb: Fast Prediction of Bacteriophage Virion Proteins Using Un-Biased Multi-Perspective Properties with Recursive Feature Elimination. \[J\] Genomics](#)
M. Arif, F Ali, S Ahmad, M Kabir, Z Ali and M Hayat;
- [2020] [iNR-2L: a two-level sequence-based predictor developed via Chou's 5-steps rule and general PseAAC for identifying nuclear receptors and their families. \[J\] Genomics.](#)
M Kabir, S Ahmad, M Iqbal, and M Hayat;

- [2020] [SDBP-Pred: Prediction of single-stranded and double- stranded DNA-binding proteins by extending consensus sequence and K-segmentation strategies into PSSM. Analytical Biochemistry.](#)
F Ali, **S Ahmad**, M Kabir, M Arif, ZU Khan, and DJ Yu;
- [2019] [DP-BINDER: machine learning model for prediction of DNA-binding proteins by fusing evolutionary and physicochemical information. \[J\] Journal of Computer-Aided Molecular Design.](#)
F Ali, **S Ahmed**, ZNK Swati, S Akbar,
- [2019] [Content-Based Brain Tumors Retrieval for MR Images Using Transfer Learning. \[J\] IEEE Access.](#)
ZNK Swati, QH Zhao, M Kabir, Z Ali, F Ali, **S Ahmad**, and JF Lu:
- [2019] [Brain Tumor Classification for MR Images using Transfer Learning and Fine-Tuning. \[J\] Computerized Medical Imaging and Graphics](#)
ZNK Swati, QH Zhao, M Kabir, **S Ahmad**, Z Ali, and JF Lu:
- [2019] [Prediction of membrane protein types by exploring local discriminative information from evolutionary profiles. \[J\] Analytical Biochemistry.](#)
M Kabir, M Arif, **S Ahmad**, F Ali, ZNK Swati, and DJ Yu;
- [2019] [iDTi-CSsmoteB: identification of drug-target interaction based on drug chemical structure and protein sequence using XGBoost with over-sampling technique SMOTE](#)
SMH Mahmud, W Chen, H Jahan, Y Liu, NI Sujan, **S Ahmed**;
- [2018] [An Integrated Feature Selection algorithm for Cancer Classification using Gene Expression Data. \[J\] Combinatorial Chemistry & High Throughput Screening](#)
S Ahmad, M Kabir, Z Ali, M Arif, DJ Yu;
- [2018] [Improving secretary proteins prediction in Mycobacterium tuberculosis using oversampling technique with combination of support vector machine. \[J\] International Journal of Data Mining and Bioinformatics](#)
S Ahmad, M Kabir, M Arif, Z Ali, F Ali, and ZNK Swati:
- [2018] [Intelligent computational method for discrimination of anticancer peptides by incorporating sequential and evolutionary profiles information. \[J\] Chemometrics and Intelligent Laboratory Systems](#)
M Kabir, M Arif, **S Ahmad**, Z Ali, ZNK Swati, and DJ Yu;
- [2018] [Improving prediction of extracellular matrix proteins using evolutionary information via a grey system model and asymmetric under-sampling technique. \[J\] Chemometrics and Intelligent Laboratory Systems](#)
M Kabir, M Iqbal, **S Ahmad**, ZNK Swati, Z Liu, and DJ Yu;

[Bi-PSSM: Position specific scoring matrix based intelligent computational model for identification of mycobacterial membrane proteins. \[J\] Journal of theoretical biology](#)

[2017]

M Khan, M Hayat, SA Khan, **S Ahmad**, N Iqbal;

[Identification of heat shock protein families and J-protein types by incorporating Dipeptide composition into Chou's general PseAAC. \[J\] Computer Methods and Programs in Biomedicine](#)

[2015]

S Ahmad, M Kabir, and M Hayat;

[2025]

[There will always be variants of uncertain significance. Analysis of VUSs](#)

Authors: Haoyang Zhang, Muhammad Kabir, Saeed Ahmed, Mauno Vihinen | **Journal Name:** NAR Genomics and Bioinformatics | **Volume, Issue and Pages:** 6,4,1qae154 | **Publisher:** Oxford University Press

[2015]

[iTIS-PseKNC: Identification of Translation Initiation site in human genes using pseudo k-tuple Nucleotides Composition. \[J\] Computers in Biology and Medicine](#)

M Kabir, M Iqbal, **S Ahmad**, and M Hayat;

CONFERENCES AND SEMINARS

Comprehensive analysis of machine learning based predictors for identifying DNase I hypersensitive sites

4th International Conference on Innovative Computing (IC2-2021), UMT, Lahore

M Rashid Rasheed, Mehwish Gill, M A Subhani, M Arif, **S Ahmed**, M Kabir*;

A Recent advancements in predicting protein phosphorylation sites using machine learning methods.

4th International Conference on Innovative Computing (IC2-2021), UMT, Lahore

A Yousaf, M Rashid Rasheed, M Arif, A Yousafzai, M Kabir, **S Ahmed***;

NETWORKS AND MEMBERSHIPS

[08/12/2023 – Current]

Editorial Board Member, BMC Bioinformatics'

Member of the organizing committee for the 4th ICIC Conference virtually held at UMT in 2021.

UMT Lahore

[12/08/2024 – 12/05/2025]

Guest Editor, at Journal of Discover Health Systems.

DRIVING LICENCE

Cars: B1 18/11/2025 – 25/11/2029

HONOURS AND AWARDS

Honors and Awards

- **Postdoctoral Fellowship**, Lund University, 2023–2024
- **Postgraduate Full Scholarship** awarded by the Chinese Government via the Chinese Scholarship Council (CSC) consisting of a free tuition fee, medical insurance, and a 3500¥ stipend per month for studying Ph.D. in Computer Science and Technology at Nanjing University of Science & Technology, China (September 2016 to June 2020).

- **The International Postdoctoral Fellowship** Fund 2021 was received from the Faculty of Medical Technology, University of Mahidol, Bangkok, Thailand, to support the research work from June 1, 2021, to February 28, 2022.
- **Merit Scholarship** during M.S. Abdul Wali Khan University Mardan.
- **Merit Scholarship** at the B.S. University of Science and Technology, Bannu.

THESIS SUPERVISION/ CO-SUPERVISION

Mehwish Gill Co-Supervisor - 2022

Prediction and analysis of computational methods for the identification of enhancers and their strengths.

Roha Arif Co-Supervisor - 2023

Identification of Tumor-Homing Peptides Using Deep features combined Support vector machine...

Sameera Kanwal Supervisor - 2023

Prediction of antimicrobial peptides using feature representation learning.

Abdullah Muhammad Asghar Supervisor - 2023

Large-scale comparative review and assessment of computational methods for identification of nucleosome positioning.

Farwa Arshad Co-Supervisor - 2023

Identification of anti-diabetic peptides using stacked based ensemble learning.

Aqsa Amjad Supervisor - 2023

Analysis and identification of promoters and their strength based on a deep neural network with sequential features.

Shumaila Kanwal Co-Supervisor - 2023

Prediction of ampylation sites using a protein language model with a deep cascade random forest.

Hina Farooq Supervisor - 2023

A novel approach for accurate prediction of Cyclin Proteins using stacking ensemble based learning.

Muhammad Rehan Amjad Supervisor - 2023

Prediction of Antiviral Peptides Using Long-Short-Term-Memory (LSTM)-Based Convolution Neural Networks.

ZUBDA KHANUM Supervisor - 2023

Analysis and prediction of lysine malonylation sites by exploiting informative features in machine learning framework using principal component analysis.

Zoha Kashaf Supervisor - 2023

Cell-Specific Long Non-Coding RNA Prediction Using Deep Learning Techniques.

TRAININGS ACQUIRED

Art & Science of Teaching and Learning in Higher Education

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Outcome-Based Education (OBE) from Quality Assurance Perspective

Teaching Philosophy and Course Designed by Centre of Teaching and Learning

Faculty Capacity Building Program by Centre of Teaching and Learning

THOUGHT COURSES

Courses Taught (Undergraduate Level)

- Introduction to Information and Communication Technologies
- Introduction to Computing
- Introduction to Data Science
- Discrete Structures
- Programming Fundamentals (with C++ and Python)
- Data Structure and Algorithm
- Machine Learning
- Deep Learning
- Data Structures and Algorithms

MANAGEMENT AND LEADERSHIP SKILLS

Batch Advisor for the student of BS-Computer Science and PhD-Computer Science.

Providing counsel and support to students as their adviser, assisting them with academic and career-related decisions.

NCEAC Committee

Monitored and prepared all activities for the NCEAC visit for accreditation of BS-CS degrees at the CS Department and ensured a successful visit.

Co-curricular, extra-curricular, and external events Committee

Enhance and manage both the co-curricular and extra-curricular activities at school, including organizing external events

Conference organizing committee

Collaborating with other committee members to plan and organize all aspects of the conference, including setting objectives, defining themes, and determining the conference schedule,.

[01/10/2024 – Current]

BS coordinator of Software Engineering

REFEREE SERVICES

Referee services

- Briefings in Bioinformatics
- Scientific Reports
- IEEE Access.
- Genomics.
- Artificial Intelligence in Medicine.
- Computers in Biology and Medicine.
- Visual Computing for Industry, Biomedicine, and Art

- Genes (MDPI)
- Electronic (MDPI)
- International Journal of Molecular Science (MDPI)
- International Conference on Innovative Computing – UMT
- International Conference on Frontiers of Information Technology – COMSATS
- Journal of Soft Computing
- Applied Soft computing
- Advance Bioinformatics (Oxford press)
- PLOS One

REFERENCES

[10/09/2023]

Reference-1

Name: Prof. Dr. Mauno Vihinen

Position: Full Professor

Contact: +46725260022

Email: mauno.vihinen@med.lu.se

Institute: Faculty of Medicine, Department of Experimental Medical Science

Webpage: <https://portal.research.lu.se/en/persons/mauno-vihinen>

References-2

Name: Dr. Watshara Shoombuatong

Position: Associate Professor

Contact: +66 939384256

Email: watshara.sho@mahidol.ac.th

Institute: Center of Data Mining and Biomedical Informatics, University of Mahidol Thailand, Bangkok.

Reference-3

Name: Prof. Dong-Jun Yu

Position: Professor

Contact:

Email: njyudj@njust.edu.cn

Mobile No. +86-136-01456676

Office No. +86-25-84315751

Institute: School of Computer Science and Engineering, Nanjing University of Science & Technology, P.R. China.

Webpage: <http://csbio.njust.edu.cn/djyu/>

Reference-4

Name: Dr. Maqsood Hayat

Position: Associate Professor

Contact: +92-308-5598123

Email: m.hayat@awkum.edu.pk or Maqsood.hayat@gmail.com

Institute: Department of Computer Science, Abdul Wali Khan University Mardan, Khyber Pakhtunkhwa, Pakistan.

Webpage: https://www.awkum.edu.pk/Departments/Computer_Science/CS_Faculty_Maqsood_Hayat.html