Md Hossain Shuvo, Ph.D.

Assistant Professor

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EDUCATION

DEC 2023 Virginia Tech

Ph.D. in Computer Science

Advisor: Dr. Debswapna Bhattacharya

GPA: 3.88/4.0

JUL 2017 Alabama A&M University

M.Sc. in Computer Science

Advisor: Dr. Yujian Fu

GPA: 4.0/4.0

JUL 2014 Bangladesh University of Business and Technology

B.Sc. in Computer Science

Advisor: Mr. Md Saifur Rahman

CGPA: 3.86/4.0

RESEARCH INTERESTS

Computational Biology
Machine Learning

Data Analytics

Artificial Intelligence

RESEARCH EXPERIENCE

AUG 2021 – DEC 2023 Virginia Tech

Advisor: Dr. Debswapna Bhattacharya

Research direction: application of machine learning in developing computational methods for addressing problems

related to protein complexes

JAN 2018 – JUL 2021 Auburn University

Advisor: Dr. Debswapna Bhattacharya

Research direction: application of machine learning in developing computational methods for addressing problems

related to protein structure prediction

AUG 2015 – JUL 2017 Alabama A&M University

Advisor: Dr. Yujian Fu

Research direction: development of tools for analyzing interactive behaviors in embedded and heterogeneous robotic

systems

ACADEMIC EXPERIENCE

JAN 2024 - PRESENT Prairie View A&M University

Position: Assistant Professor, Dept. of CS

AUG 2018 – JAN 2020 Auburn University

Position: Graduate Teaching Assistant, Dept of CSSE

Courses:

COMP 5970/6970: Computational Biology
 COMP1210: Fundamental of Computing I

MAY 2016 – JUL 2016 North Alabama Center for Educational Excellence (NACEE)

Position: Assistant Mentor

Topic: implementing multi-tasking and interactive behavior in

NAO Humanoid robot

JAN 2015 – JUL 2015 Dhaka Commerce College, Dhaka, Bangladesh

Position: Lecturer, Dept. of Stat, Math and COMP

JAN 2014 – DEC 2014 Dhaka Cambrian College, Dhaka, Bangladesh

Position: Lecturer, Dept. of Information and Communication

Technology

REFEREED PUBLICATIONS

2025

18. **M. H. Shuvo**, D. Bhattacharya. EquiRank: improved protein-protein interface quality estimation using protein-language-model-informed equivariant graph neural networks International Conference on Intelligent Biology and Medicine, 2024 Computational and Structural Biotechnology Journal, Volume 27, 160 - 170, 2025

2024

17. R. Roche, B. Moussad, **M. H. Shuvo**, S. Tarafder, D. Bhattacharya, EquiPNAS: improved protein-nucleic acid binding site prediction using protein-language-model-informed equivariant deep graph neural networks. *Nucleic Acids Research*, 2024 gkae039, 10.1093/nar/gkae039

2023

- 16. **M. H. Shuvo**, M. Karim, R. Roche, and D. Bhattacharya, "PIQLE: protein-protein interface quality estimation by deep graph learning of multimeric interaction geometries", Bioinformatics Advances, 2023, vbad070, 10.1093/bioadv070.
- 15. **M. H. Shuvo**, M. Karim, and D. Bhattacharya, "iQDeep: an integrated web server for protein scoring using multiscale deep learning models", Journal of Molecular Biology, 168057, 2023. doi: 10.1016/j.jmb.2023.168057
- 14. R. Roche, B. Moussad, **M. H. Shuvo**, D. Bhattacharya, "E(3) equivariant graph neural networks for robust and accurate protein–protein interaction site prediction", PLOS Computational Biology, 19, e1011435, doi: 10.1371/journal.pcbi.1011435
- 13. S. Bhattacharya, R. Roche, **M. H. Shuvo**, and D. Bhattacharya, "Contact-assisted threading in low-homology protein modeling", Methods in Molecular Biology book series, vol. 2627, 2023, doi: 10.1007/978-1-0716-2974-1 3

2022

12. R. Roche, S. Bhattacharya, **M. H. Shuvo**, and D. Bhattacharya, "rrQNet: Protein contact map quality estimation by deep evolutionary reconciliation", Proteins, Jun 2022, doi: 10.1002/prot.26394.

2021

- M. H. Shuvo, M. Gulfam, and D. Bhattacharya, "DeepRefiner: high-accuracy protein structure refinement by deep network calibration", Nucleic Acids Research, vol. 49, no. W1, pp. W147– W152, Jul. 2021, doi: 10.1093/nar/gkab361
- 10. S. Bhattacharya, R. Roche, **M. H. Shuvo**, and D. Bhattacharya, "Recent Advances in Protein Homology Detection Propelled by Inter-Residue Interaction Map Threading", Front Mol Biosci, vol. 8, p. 643752, 2021, doi: 10.3389/fmolb.2021.643752.
- 9. A. Kryshtafovych, . . . , **M. H. Shuvo**, . . . , "Modeling SARS-CoV-2 proteins in the CASP-commons experiment", Proteins, vol. 89, no. 12, pp. 1987–1996, Dec. 2021, doi: 10.1002/prot.26231.

2020

- 8. **M. H. Shuvo**, S. Bhattacharya, and D. Bhattacharya, "QDeep: distance-based protein model quality estimation by residue-level ensemble error classifications using stacked deep residual neural networks", Bioinformatics, vol. 36, no. Suppl 1, pp. i285–i291, Jul. 2020, doi: 10.1093/bioinformatics/btaa455
- 7. R. Alapati, **M. H. Shuvo**, and D. Bhattacharya, "SPECS: Integration of side-chain orientation and global distance-based measures for improved evaluation of protein structural models", PLoS One, vol. 15, no. 2, p. e0228245, 2020, doi: 10.1371/journal.pone.0228245

2016

- 6. **M.H. Shuvo**, Y. Fu, "Sonar sensor virtualization for object detection and localization", SoutheastCon 2016, 1–8. doi: 10.1109/SECON.2016.7506699
- 5. Y. Fu, **M.H. Shuvo**, "An approach to analyzing adaptive intelligent vehicle system using SMT solver", In 2016 International Conference on Control, Decision and Information Technologies", (CoDIT)., pp. 313–319, doi: 10.1109/CoDIT.2016.7593580
- 4. Y. Fu, **M.H. Shuvo**, "Android-Based Remote Robot Control System". In, 2016 ASEE Annual Conference & Exposition Proceedings. ASEE Conferences, New Orleans, Louisiana, p.26251, doi: 10.18260/p.26251

2015

- 3. S. Rahman, **M.H. Shuvo**, "Advancement of Information System in the Health Sphere in Rural-Urban Areas of Developing Countries: A Case Study of Bangladesh". IJIRSET, Vol. 4, Issue 6, June 2015, doi: 10.15680/IJIRSET.2015.0406131
- 2. **M.H. Shuvo,** T. Haque, "Suspicious Behavior Detection Framework for Social Networking Sites Using Hidden Markov Model", 2015 Dhaka Commerce College Journal, Vol 7, Issue 1, P. 173 192

2014

1. M.M. Rahman, S. Ahmed, **M.H. Shuvo**, "Nearest Neighbor Classifier Method for Making Loan Decision in Commercial Bank". IJISA, 6, 60–68, doi: 10.5815/ijisa.2014.08.07

ABSTRACTS AND POSTERS

Abstracts

- 9. **M. H. Shuvo**, M. Karim, and D. Bhattacharya, "Protein modeling and accuracy estimation by Bhattacharya group in CASP15," CASP15 abstract, p. 35, 2022.
- 8. **M. H. Shuvo**, M. Gulfam, and D. Bhattacharya, "Deep network calibration for protein structure refinement," 13th ACM Conference on Bioinformatics, Computational Biology, and Health Informatics, ACM-BCB 2022.
- 7. **M. H. Shuvo**, S. Bhattacharya, R. Roche, and D. Bhattacharya, "Protein tertiary structure prediction by Bhattacharya group in CASP14," CASP14 abstract, p. 38, 2020.
- 6. **M. H. Shuvo**, and D. Bhattacharya, "Protein model accuracy estimation by Bhattacharya groups in CASP14," CASP14 abstract, p. 39, 2020.
- 5. **M. H. Shuvo**, and D. Bhattacharya, "Protein structure refinement by Bhattacharya groups in CASP14," CASP14 abstract, pp. 40–41, 2020.
- 4. D. Bhattacharya, R. Alapati, and **M. H. Shuvo**, "Protein structure prediction and refinement by Bhattacharya human group in CASP13," CASP13 abstract, pp. 29-30, 2018.
- 3. R. Alapati, **M. H. Shuvo**, and D. Bhattacharya, "clustQ: Multi-model QA using superposition-free weighted internal distance comparisons," CASP13 abstract, p. 31, 2018
- 2. D. Bhattacharya and **M. H. Shuvo**, "refineD: Protein structure refinement using machine learning guided restrained relaxation," CASP13 abstract, p. 32, 2018
- 1. D. Bhattacharya and **M. H. Shuvo**, "scoreD: Estimating Global Distance Test using deep discriminative binary classifier ensemble," CASP13 abstract, p. 33

Posters

- 3. **M. H. Shuvo**, D. Bhattacharya. "EquiRank: improved protein-protein interface quality estimation using protein-language-model-informed equivariant graph neural networks", PVAMU R&I Week, 2024
- S. Adepoju, M.H. Shuvo, "How Machine Learning Can Revolutionize Building Comfort: Unveiling the Impact of Occupancy Prediction Models on HVAC Control System of the Roy G. Perry College of Engineering Building", PVAMU R&I Week, 2024
- M. H. Shuvo, S. Bhattacharya, D. Bhattacharya, "QDeep: distance-based protein model quality estimation by residue-level ensemble error classifications using stacked deep residual neural networks", ISMB 2020

PARTICIPATION IN COMMUNITY-WIDE ASSESSMENT

APR 2022 – AUG 2022 15th Critical Assessment of Protein Structure Prediction

(CASP15) challenge

Role: method development, technical support for double-blind testing of our developed methods in both server and human

pipelines

MAY 2020 – SEP 2020 14th Critical Assessment of Protein Structure Prediction

(CASP14) challenge

Role: method development, technical support for double-blind testing of our developed methods in both server and human

pipelines

APR 2020 CASP Commons (COVID-19, 2020), a collaborative initiative

for modeling the SARS-2-CoV structure

Role: Provided technical support for testing our developed methods in predicting and assessing SARS-2-CoV target

MAY 2018 – AUG 2018 13th Critical Assessment of Protein Structure Prediction

(CASP13) challenge

Role: method development, technical support for double-blind testing of our developed methods in both server and human

pipelines

HONORS AND AWARDS

Quantum Workshop Travel

Fellowship

By the University of Tennessee at Chattanooga

Google Cloud Research Resource

\$5,000 resource credit by Google

The PHIT Workforce Development

Program Grant [Co-PI]

\$432,078 by the University of Texas Health Science Center at Houston

GenAl Faculty Summit Travel

Fellowship

Awarded by Google

FACULTY ENHANCEMENT

PROGRAM

Awarded by the Roy G. Perry College of Engineering at PVAMU

NSF NDSA PFx instructor

Selected to be a PFx instructor

NSF NDSA RESEARCH AFFINITY COHORT

Selected as 1 of the 20 researchers

GOOGLE TEC EQUITY IMPACT

FUND

\$50,000 non-dilutive funding, led jointly with Dr. Wang at PVAMU

ACCESS ALLOCATIONS

Principal Investigator (PI), NSF - XSEDE Research Allocation

PRATT FELLOWSHIP AWARD

Awarded Pratt Fellowship at Virginia Tech, 2023

YOUNG SCIENTIST EXCELLENCE

AWARDS

Awarded 1st place prize at 18th annual MCBIOS conference, 2022

CONFERENCE FELLOWSHIP

I. Received fellowship award for MCBIOS 2022

II. Received ISMB 2020 fellowship award

PUBLICATION RECOGNITION DeepRefiner paper accepted for ACM-BCB 2022 Highlights track

TRAVEL FELLOWSHIP Received travel grant for IEEE SoutheastCon 2016 Awarded 2nd place prize at AAMU STEM Day 2016 **POSTER AWARD**

SCIENTIFIC SOFTWARE DEVELOPMENT

pLM-based Protein-protein interface quality estimation method **EquiRank**

[GitHub]

PIQLE Protein-protein interface quality estimation method [GitHub]

iQDeep Integrated protein scoring server [Server]

DeepRefiner High-accuracy protein structure refinement server [Server] Single-model protein quality estimation method [GitHub] **QDeep SPECS** Improved evaluation method for protein structures [Github]

EquiPPIS E(3) equivariant graph neural networks for robust and accurate

protein-protein interaction site prediction [Github]

EquiPNAS EquiPNAS: improved protein-nucleic acid binding site prediction

using protein-language-model-informed equivariant deep

graph neural networks [GitHub]

rrQNet Protein contact map evaluation method [Github]

TECHNICAL SKILLS

PROGRAMMING SKILLS Python, R, Java, C, C++, PHP, ASP, Apache Cordova MySQL,

MSSQL

APP DEVELOPMENT I. Standalone application development with Java EE

II. Mobile applications development in both Android and iOS

III. Platforms Web applications development

IV. Robotic applications development for EV3, NAO, and UAV

SERVICES AND OUTREACH

COMPUTER SCIENCE CURRICULUM Serving as a member of the CS Undergraduate Curriculum

DEVELOPMENT COMMITTEE Development Committee

COLLEGE OF ENGINEERING CURRICULUM DEVELOPMENT

COMMITTEE

DIVISION OF RESEARCH AND

Served as a member of the COE Undergraduate Curriculum

Development Committee

INNOVATION COMMITTEE

Serving as a member of the Division of Research and Innovation

Committee for Computer Science at PVAMU

FACULTY SEARCH COMMITTEE

Served as a member of the PVAMU faculty search committee Served as a sub-reviewer for Bioinformatics, BIOKDD 2021,

REVIEWER

2023, 2024, IEEE ACM/Transaction, IEEE CogMI and others

EVENT MANAGEMENT

- i. Provided logistic support on E-day 2020 at Auburn University ii. Provided logistic support on AAMU Senior High School Day 2016

TECHNICAL COMMITTEE MEMBER Provided technical support by developing the complete submission management system for AAMU STEM Day 2016