

Syed Ashiqur Rahman, Post-doctoral Researcher, University of Pittsburgh

- Contact: sar210@pitt.edu, +1(304)-276-9808 • Website: <https://syedashiqurrahman.github.io/>
- LinkedIn: <https://www.linkedin.com/in/syed-ashiqur-rahman/> • GoogleScholar: <https://scholar.google.com/citations>

EXPERIENCE	Post-doctoral Researcher at University of Pittsburgh <i>Feb 2020 - till-date</i> <ul style="list-style-type: none">• Research with Dr. Jishnu Das on Machine Learning, Bioinformatics Graduate Research Assistant at West Virginia University <i>Jan 2012 - Dec 2019</i> <ul style="list-style-type: none">• Research with Advisor Dr. Donald Adjeroh on Deep Learning, Machine Learning, Health Informatics Graduate Teaching Assistant at West Virginia University <i>Aug 2015 - Dec 2019</i> <ul style="list-style-type: none">• CS 350: Operating Systems, grading, software lab (in C++) Course Instructor, Dept. of CSE at Daffodil International University <i>Jan 2009 - Dec 2011</i> <ul style="list-style-type: none">• Taught Undergraduate Courses (Algorithm, Data Structures, Programming)• Coach of ACM Programming Team Radio Frequency Engineer, R&D-Team at Ericsson Ltd. Bangladesh <i>Jan 2008 - Jan 2009</i> <ul style="list-style-type: none">• Optimized radio network design in the full R&D cycle (field data collection to deployment & Integration)
EDUCATION	Ph.D. in Computer Science, West Virginia University <i>Graduated: December 2019</i> <ul style="list-style-type: none">• Dissertation Topic: <i>Quantifying Human Biological Age: A Machine Learning Approach</i>• Advisor: Dr. Donald Adjeroh, Professor, Department of Computer Science, West Virginia University• CGPA: 3.93 in scale of 4.00 B.Sc. in Computer Science and Engineering <i>Graduated: December 2007</i> University of Dhaka , Dhaka, Bangladesh <ul style="list-style-type: none">• CGPA: 3.87 in scale of 4.00
ML COURSES DURING PHD	Machine Learning, Advanced Data Mining, Pattern Recognition, Cyber Security & Big-Data, Deep Learning.
TECHNICAL SKILLS	Programming Languages R (v3.62), Python (v2.7, v3.6) , Java, SQL, C/C++, Shell Scripting, HTML, Javascript, Matlab APIs and libraries Keras, Tensorflow, ScanPy, Seurat, Scikit-learn, SciPy, NumPy, MySQL, Hadoop, MapReduce, SPSS Tools RStudio, PyCharm, VSCode, Adobe Illustrator, Git, L ^A T _E X, Weka, SVN, Cytoscape, Inkscape Operating Systems macOS Big Sur, Ubuntu 16.04, Redhat Linux, Windows 10
SELECTED PROJECTS	Using interpretable machine learning approaches to uncover immune signatures of Covid-19 outcome using Python, R <ul style="list-style-type: none">• Applied machine learning model, a latent factor based regression, and a causal graphical model to infer predictive performance and molecular mechanistics for ICU Covid-19 patients (cytokine and chemokine)• Showed inference beyond prediction for the ICU patients bifurcating between survivors and non-survivors Non-Spike protein and endemic coronavirus antibodies associated with outcomes in severe Covid-19 using Python, R <ul style="list-style-type: none">• Applied a machine learning model and a latent factor based regression model to infer predictive performance and molecular mechanistics for ICU Covid-19 patients using Antibody-omic profiling. Network-based integration of epigenetic landscapes unveils molecular programs underlying human T follicular helper cell (Tfh) differentiation using Python, R <ul style="list-style-type: none">• Combined epigenomic datasets and integrated them with the reference human protein interactome using a novel network propagation approach• Uncovered subnetworks integral to Tfh cell differentiation A network-based approach to identify expression modules underlying rejection in pediatric liver transplantation using Python, R <ul style="list-style-type: none">• Discovered pre and post transplant molecular signatures of liver transplant outcome from whole blood transcriptomes using machine learning• Combined transcriptomic data with high-quality human protein interactome network to identify differentially regulated functional sub-components of the network A machine learning approach to identify expression modules underlying rejection in kidney transplantation using Python, R <ul style="list-style-type: none">• Showed discrimination between clinical outcomes using kidney-compartment specific protein abundances combined with modularity of the underlying protein interactome network Anthropometric measurements to MakeHuman using Python, R, Keras, Tensorflow <ul style="list-style-type: none">• Built a plugin to generate MakeHuman 3D models from NHANES body measurements• Generated 3D models are now being used for health profiling applying Deep Learning techniques

Mini-places Challenge : Scene Recognition by CNN using Python, Keras, Tensorflow

- Used MiniPlaces dataset for scene recognition using Deep CNN
- Achieved improved performance in compare to baseline refNet1

Vehicle Brand Identification using Matlab, SIFT, SURF, SVM

- Images were collected from west virginia university (WVU) parking lot. This is part of an intelligent traffic system where statistics of the various car passing in a given zone is of interest.

Surface-based Body Shape Index (SBSI) and its relation with all-cause mortality using R, Survival Statistical Models (CoxPH, Kaplan-Meier)

- Introduced a new body measurement index called SBSI
- SBSI is linear with age, and increases with increasing mortality, performs better than BMI

Convolutional LSTM estimates biological age from human physical activity using R, Keras

- Introduced an innovative 3D architecture to find patterns in human physical activity
- Proposed 3D approach beats the 1D and 2D deep learning based methods

SELECTED PUBLICATIONS

Journal

- VS Mahajan, **SA Rahman**, VV Viswanadham, GJ Yuen, N Sun, H Mattoo, SS Pillai, J Das, “[Network-based integration of epigenetic landscapes unveils molecular programs underlying human T follicular helper cell differentiation](#)”, bioRxiv, 2021 (under review in Cell Reports).
- **SA Rahman**, P Giacobbi, L Pyles, C Mullett, G Doretto, D Adjeroh, “[Deep learning for biological age estimation](#)”, Briefings in Bioinformatics, 2020.
- **SA Rahman**, D Adjeroh, “[Deep Learning using Convolutional LSTM estimates Biological Age from Locomotor Physical Activity](#)”, Nature Scientific Reports, 2019.
- **SA Rahman**, D Adjeroh, “[Centroid of Age Neighborhoods: A New Approach to Estimate Biological Age](#)”, Journal of Biomedical and Health Informatics, 2019.
- **SA Rahman**, D Adjeroh, “[Surface-Based Body Shape Index and Its Relationship with All-Cause Mortality](#)”, PLoS ONE 10 (12), 2015.

Conference

- **SA Rahman**, D Adjeroh, “Estimating Biological Age from Physical Activity using Deep Learning with 3D CNN”, [IEEE International Conference on Bioinformatics and Biomedicine](#), 2019.
- **SA Rahman**, D Adjeroh, “Centroid of Age Neighborhoods: A Generalized Approach to Estimate Biological Age”, [The IEEE-EMBS International Conference on Biomedical and Health Informatics](#), 2019.

HONORS & AWARDS

News

- Our collaborative research received \$4 million grant from National Science Foundation and was highlighted in WVU’s College of Engineering. [<https://www.statler.wvu.edu/news/2019/10/09/>]

Professional Services

- SBP-BRiMS Program Committee, 2019
- SBP-BRiMS Student Chair, 2015

Travel Grants

- IEEE BHI Student Travel Award (NSF), 2019
- CITeR Student Travel Award, 2018
- SBP-BRiMS Student Travel Award, 2016, 2014

Merit Scholarships

- Full Tuition Waiver Scholarship, B.Sc. program in Computer Science, University of Dhaka
- Ministry of Education, Government of Bangladesh
 - Excellence in Higher Secondary Certificate Exam, 2001-2005
 - Excellence in Secondary School Certificate Exam, 1999-2001

Extra Curricular Activities

- WVU Intramural Pingpong Champion, 2015
- Chess Champion in High School

REFERENCE

Available upon request.