# Mohammad Shabbir Hasan

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Career OBJECTIVE

To be a successful researcher in Computer Science and build up a career as an academician.

Research Interests Computational Biology; Bioinformatics; Variant Calling; Evolutionary Genomics.

EDUCATION

Virginia Tech, USA.

Ph.D. in Computer Science,

Expected: Summer 2018

- Thesis Topic: Identifying genetic variations in human genome.
- Advisor: Liqing Zhang, Ph.D., Associate Professor, Department of Computer Science, Virginia Tech.

### The University of Akron, USA.

Masters in Computer Science,

using clustering algorithms.

• Thesis Topic: Investigating gene relationships in microarray expressions: approaches

Graduated: August 2013

• Advisor: Zhong-Hui Duan, Ph.D., Professor, Department of Computer Science, The University of Akron.

#### Khulna University of Engineering and Technology, Bangladesh.

B.Sc.(Engr.) in Computer Science and Engineering, Graduated: March 2008

- Thesis Topic: A study on reducing the development cost for reusable Object Oriented software.
- Advisor: K.M. Azharul Hasan, Ph.D., Professor, Department of Computer Science and Engineering, Khulna University of Engineering and Technology.

### Work Experience

#### Research Assistant

August 2013 to present

• Computational Biology and Bioinformatics Lab (Zhang Lab), Virginia Tech

Research Projects:

- Genesis-indel: Gene set Influenced Analysis of indels.
- UPS-indel: A Universal Positioning System for indels.
- P-Dindel: A multi-thread based tool for calling indels from short reads.
- SPAI: Single Platform for Analyzing Indels.
- Performance evaluation of indel calling tools using real short-read data.

### Software Development Engineering Intern

May 2016 to August 2016

• Amazon.

• Development of a Coral service application for the Amazon Prime customers.

## Software Engineering Intern

May 2015 to August 2015

• Biotronik/Micro Systems Engineering Inc.

#### Projects:

• Improved communication for implants using command batching.

#### Research Assistant

August 2011 to July 2013

• Department of Computer Science, The University of Akron.

#### Research Projects:

• Hierarchical k-Means: A hybrid clustering algorithm and its application to study gene expression in Lung Adenocarcinoma.

#### Research Assistant

January 2011 to July 2011

• Che Lab of BigData, East Stroudsburg University of Pennsylvania.

#### Research Projects:

- EGID: An Ensemble algorithm for Genomic Island Detection.
- GIST: Genomic Island Suite of Tools.

## TEACHING EXPERIENCE

#### Teaching Assistant

August 2013 to present

• Department of Computer Science, Virginia Tech

#### Responsibilities:

• Teaching Lab classes for the courses Introduction to Programming in Java, Introduction to Software Design, and Intermediate Software Design.

#### Teaching Assistant

August 2011 to July 2013

• Department of Computer Science, The University of Akron.

#### Responsibilities:

• Assisting professors for the courses Algorithm, Data Structures, Object Oriented Programming, and Advanced Algorithm.

#### Lecturer

January 2010 to December 2010

• Department of Computer Science, Institute of Science, Trade, and Technology, Dhaka, Bangladesh.

### Responsibilities:

- Teaching theory and lab classes on **Data Structures**, **Microprocessors** for undergraduate classes.
- Grading exams, projects, and assignments.

## Lecturer

May 2008 to September 2009

• Department of Computer Science, Asian University of Bangladesh, Dhaka, Bangladesh.

## Responsibilities:

- Teaching theory and lab classes on Algorithms, Discrete Mathematics, Introduction to Programming in C, Internet Programming for undergraduate classes.
- Grading exams, projects, and assignments.

### BOOK CHAPTER

1. **Hasan, M.S.** and Duan, Zhong-Hui. "Hierarchical k-Means: A hybrid clustering algorithm and its application to study gene expression in Lung Adenocarcinoma." in "Emerging Trends in Computational Biology, Bioinformatics, and Systems Biology - Algorithms and Software Tools.", Elsevier, pp. 51-67, 2015.

### Journal Publications

- Hasan, K.M.A. and Hasan, M.S. "A parsing scheme for finding the design pattern and reducing the development cost of reusable object oriented software." International Journal on Computer Science and Information Technology, 2(3):40– 54, 2010.
- 2. **Hasan, M.S.**, Farhan, Q., and Al Mahmood, A. "An exploratory and feasibility study of implementing online based voting system in Bangladesh." *International Journal of Computer Science and Emerging Technologies*, 1(3):125–132, 2010.
- 3. Hasan, M.S., Al Mahmood, A., Alam, Md., Hasan, Sk.Md., and Rahman, F. "An evaluation of software requirement prioritization techniques." *International Journal of Computer Science and Information Security*, 8(9):83–94, 2010.
- Hasan, Sk.Md., Hasan, M.S., Al Mahmood, A., and Alam, Md. "A model for value based requirement engineering." *International Journal of Computer Science and Network Security*, 10(12):171–177, 2010.
- 5. Che, D., **Hasan, M.S.**, Wang, H., Fazekas, J., Huang, J., and Liu, Q. "EGID: an ensemble algorithm for improved genomic island detection in genomic sequences." *Bioinformation*, 7(6):311–314, 2011.
- Hasan, M.S., Liu, Q., Wang, H., Fazekas, J., Chen, B., and Che, D. "GIST: Genomic Island Suite of Tools for predicting genomic islands in genomic sequences." *Bioinformation*, 8(4):203–205, 2012.
- 7. Che, D., **Hasan, M.S.**, and Chen, B. "Identifying pathogenicity islands in bacterial pathogenomics using computational approaches." *Pathogens*, 3(1):36–56, 2014.
- 8. **Hasan, M.S.**, Wu,X., and Zhang, L. "Performance evaluation of indel calling tools using real short-read data." *Human Genomics*, 9(1):1–14, 2015.
- 9. **Hasan, M.S.**, and Zhang, L. "SPAI: an interactive platform for indel analysis." *BMC Genomics*, 17(5):496, 2016.
- 10. **Hasan, M.S.**, Wu, X., Watson, L., and Zhang, L. "UPS-indel: A Universal Positioning System for indels." *Scientific Reports*, 7(1), 2017

## Conference Publications

- 1. **Hasan, M.S.** and Hasan, K.M.A. "Finding the design pattern from the source code for developing reusable object oriented software."  $2^{nd}$  International Conference on the Applications of Digital Information and Web Technologies, pp.157–162, London, UK, August 2009.
- 2. Hasan, K.M.A. and **Hasan, M.S.** "Principal component analysis of coupling measures for developing high quality object oriented software." 3<sup>rd</sup> International Conference on Computer and Communication Engineering, pp.217–222, Kuala Lumpur, Malaysia, July 2010.
- 3. Hasan, M.S. and Hasan, K.M.A. "Determining the Most Effective Class for Extending Reusability of Object Oriented System." *International Conference on Educational and Information Technology*, pp. Volume 2: 498–502, Chongqing, China, September 2010.
- 4. **Hasan, M.S.**, Al Mahmood, A., and Farhan, Q. "A roadmap towards the implementation of an efficient online voting system in Bangladesh." *International Conference on Software Engineering and Computational Intelligence*, pp. 1–4, Wuhan, China, December 2010.
- 5. Che, D., **Hasan, M.S.**, Wang, H., Chen, B., and Wei, Y. "M are better than one: An ensemble method for genomic island prediction." 6<sup>th</sup> International Conference on Bioinformatics and Biomedical Engineering, pp.Volume 2:426–429, Shanghai, China, May 2012.

 Hasan, M.S. and Duan, Z. "A Hybrid Clustering Algorithms and Functional Study of Gene Expression in Lung Adenocarcinoma." World Comp: International Conference on Bioinformatics and Computational Biology, pp. 23–29, Las Vegas, USA, July 2014.

#### SHORT PAPERS

- 1. **Hasan, M.S.** and Zhang, L. "SPAI: Single Platform for Analyzing Indels." 11<sup>th</sup> International Symposium on Bioinformatics Research and Applications (ISBRA), pp.75–78, Norfolk, USA, June 2015.
- 2. **Hasan, M.S.** and Zhang, L. "P-Dindel: A multi-thread based tool for calling indels from short reads." 11<sup>th</sup> International Symposium on Bioinformatics Research and Applications (ISBRA), pp.71–74, Norfolk, USA, June 2015.
- 3. Tithi, S. and **Hasan, M.S.** "Modeling ebola outbreak: a case study on 2014 outbreak in sierra leone." 6<sup>th</sup> ACM International Conference on Bioinformatics, Computational Biology and Biomedicine (ACM BCB), pp.547–548, Atlanta, USA, September 2015.
- 4. **Hasan, M.S.**, Tithi, S., Tilevich, E., and Zhang, L. "Diagnosing and Treating Code-Duplication Problems in Bioinformatics Libraries." 6<sup>th</sup> IEEE International Conference on Computational Advances in Bio and Medical Sciences (ICCABS), Atlanta, USA, October 2016.
- 5. **Hasan, M.S.**, Wu, X., Watson, L., Li, Z., and Zhang, L. "UPS-indel: A better approach for finding indel redundancy." 6<sup>th</sup> IEEE International Conference on Computational Advances in Bio and Medical Sciences (ICCABS), Atlanta, USA, October 2016.

## VOLUNTARY ACTIVITIES

#### Reviewer

2014 to present

- Scientific Reports (publisher: Nature)
- Briefings in Bioinformatics (publisher: Oxford University Press)
- American Journal of Bioinformatics and Computational Biology

## Honors and Awards

#### Research Award

2017

• Microsoft Azure Research Award.

Travel Grant 2013 to present

- NSF Travel Fellowship to attend IEEE ICCABS 2016.
- Virginia Tech Graduate Student Assembly Travel Fellowship to attend IEEE ICCABS 2016.
- NSF Travel Fellowship to attend ISBRA 2015.
- Sweden-Bangladesh Trust Fund.

#### FDR Fellowship

2011

• East Stroudsburg University of Pennsylvania.

#### Merit Scholarship

2005 to 2008

• Khulna University of Engineering and Technology, Bangladesh.

RECENT TALKS

• "UPS-indel: a Universal Positioning System for Indels",  $32^{nd}$  Graduate Student Assembly (GSA) Research Symposium, Virginia Tech, March 23, 2016.

### References

Available upon request