HUSSEIN MOHSEN

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EDUCATION

Yale University, New Haven, CT, USA

Doctor of Philosophy (PhD) in Computational Biology & Bioinformatics

Aug 2016-present

Committee: Mark Gerstein, Lajos Pusztai, Kei-Hoi Cheung, Sahand Negahban Thesis Title: Network Approaches to the Study of Genomic Variation in Cancer

Master of Arts (MA) in History of Science & Medicine

Jan-Dec 2019

Graduate Certificate in Public Humanities

Jan 2021-present

Certificate Capstone Project: A Pedagogical History of the "Population" Concept in Genetics from Mid-19th Century to Current Ancestry Tests

Indiana University, Bloomington, IN, USA

Aug 2013-May 2015

Master of Science (MS) in Computer Science (Track: Bioinformatics)

Newcastle University, Newcastle upon Tyne, UK

Sep 2012-Aug 2013

Postgraduate Exchange Student at the School of Computing Science

Lebanese American University, Beirut, Lebanon

Sep 2008-Jun 2011

Bachelor of Science (BS) in Computer Science with High Distinction

PUBLICATIONS

Research Papers

- **H. Mohsen**, V. Gunasekharan, T. Qing, M. Seay, Y. Surovtseva, S. Negahban, Z. Szallasi, L. Pusztai, and M. Gerstein (2021). Network propagation-based prioritization of long tail genes in 17 cancer types, *Genome Biology*, 22, 287.
- T. Qing, **H. Mohsen**, M. Marczyk, Y. Ye, T. O'Meara, H. Zhao, J.P. Townsend, M. Gerstein, C. Hatzis, Y. Kluger and L. Pusztai (2020). Germline variant burden in cancer genes correlates with age at diagnosis and somatic mutation burden, *Nature Communications*, 11, 2438.
- **H. Mohsen**, J. Warrell, M.R. Min, S. Negahban, and M. Gerstein (2020). Weight-based Neural Network Interpretability using Activation Tuning and Personalized Products, *Machine Learning in Computational Biology Workshop (MLCB'20)*.

- M. Amodio, D. van Dijk, K. Srinivasan, W.S. Chen, **H. Mohsen**, K.R. Moon, A. Campbell, Y. Zhao, X. Wang, M. Venkataswamy, A. Desai, V. Ravi, P. Kumar, R. Montgomery, G. Wolf, and S. Krishnaswamy (2019). Exploring Single-Cell Data with Deep Multitasking Neural Networks, *Nature Methods*, 16, pp. 1139–1145.
- S. Lou, K.A. Cotter, T. Li, J. Liang, **H. Mohsen**, J. Liu, J. Zhang, S. Cohen, J. Xu, H. Yu, M. Rubin, and M. Gerstein (2019). GRAM: A generalized model to predict the molecular effect of a non-coding variant in a cell-type specific manner, *PLoS Genetics*, 15 (8): e1007860.
- J. Warrell, **H. Mohsen**, and M. Gerstein (2018). Rank Projection Trees for Multilevel Neural Network Interpretation, *NeurIPS Machine Learning for Health Workshop (NeurIPS'18 ML4H)*.
- **H. Mohsen**, H. Tang, and Y. Ye (2017). DNPipe: Improving De Novo Metatranscriptome Assembly via Machine Learning Algorithms, *International Journal of Computational Biology and Drug Design* (*IJCBDD*), 2 (10), pp. 91-107.
- **H. Mohsen**, H. Kurban, K. Zimmer, M. Jenne, and M. Dalkilic (2015). Red-RF: Reduced Random Forests using priority voting dynamic data reduction, *IEEE International Congress on Big Data (IEEE BigData Congress'15)*, pp. 118-125.
- **H. Mohsen**, H. Kurban, M. Jenne, and M. Dalkilic (2014). A New Set of Random Forests with Varying Dynamic Data Reduction and Voting Techniques, *IEEE International Conference on Data Science and Advanced Analytics (IEEE DSAA'14)*, pp. 309-405.
- N. Mansour and **H. Mohsen** (2014). Computational Evaluation of Protein Energy Functions, *International Conference on Intelligent Computing (ICIC'14), Lecture Notes in Computer Science (LNCS): Intelligent Computing in Bioinformatics*, 8590, pp. 288-299.
- **H. Mohsen** (2014). A Model to Measure Inter-communication between Segregated Communities, *IEEE International Conference on Behavioral, Economic and Social Computing (IEEE BESC'14)*, pp. 1-6.

Under Review

- J. Warrell, **H. Mohsen**, and M. Gerstein. Compression-based Network Interpretability Schemes, *bioRxiv*: 358226.
- J. Warrell*, H. Mohsen*, and M. Gerstein. Interpretability and Implicit Model Semantics in Biomedicine and Deep Learning.
- T. Qing*, **H. Mohsen***, V.L. Cannataro, M. Marczyk, M. Rozenblit, J. Foldi, M.F. Murray, J.P. Townsend, Y. Kluger, M. Gerstein, and L. Pusztai. Cancer Relevance of Human Genes, *bioRxiv*: 429823.

Reviews and Commentary

H. Mohsen (2020). Race and Genetics: Somber History, Troubled Present, *Yale Journal of Biology and Medicine*, 93 (1), pp. 215-219.

F.C.P. Navarro, **H. Mohsen**, C. Yan, S. Li, M. Gu, W. Meyerson, and M. Gerstein (2019). Genomics and data science: an application within an umbrella, *Genome Biology*, 20 (109).

WORK EXPERIENCE

New Books Network *History of Science* Podcast – *Host*Yale University, New Haven, CT, USA – *Teaching Fellow*Jan-May 2019, Jan-May 2020

Courses: Biomedical Data Science (Spring'19 and '20)

Lattice Engines, Inc., San Mateo, CA, USA - Research Engineer

Jun 2015-Jul 2016

Machine Learning R&D on imputation in sparse datasets

Indiana University, Bloomington, IN, USA - Associate Instructor

Aug 2013-May 2015

Courses: Data Mining (Spring'15) and Discrete Mathematics (Fall'13 and Spring '14)

CCT International, Beirut, Lebanon - Software Developer

Mar 2011-Aug 2012

SKILLS

Programming: Python, R, TensorFlow, Java, MPI **Web Development**: Javascript, PHP, HTML, CSS

Other: UNIX, Hadoop, Adobe Photoshop and Illustrator

AWARDS & HONORS

| American Association for Cancer Research (AACR) Scholar-in-Training Award | Oct 2020 |
|---|-----------|
| Franke Fellowship in Science and the Humanities | 2019-2020 |
| Gruber Science Fellowship | 2016-2019 |
| Fulbright Scholarship | 2013-2015 |
| Erasmus Mundus Scholarship | 2012-2013 |
| Best Computer Science Capstone Project Award at LAU (class of 2011) | Jul 2011 |
| 2 nd rank, Nokia-NNA contest for mobile application development in Lebanon | Jul 2010 |
| Extreme Programmer Award, ACM LCPC Contest, Beirut, Lebanon | Jul 2010 |
| Lebanese American University Honor List | 2009-2011 |
| Lebanese American University Merit Scholarship | 2008-2011 |

REFERENCES

Available upon request