
CURRENT POSITION

2018-present Faculty member, Vector Institute, Toronto, Canada
2015-present Assistant Professor, University of British Columbia (UBC), Department of Statistics, Department of Medical Genetics, Vancouver, Canada
2015-present Scientist, Center for Molecular Medicine and Therapeutics at UBC

Honors

2018-present Canada CIFAR AI Chair
2015-present Canada Research Chair (CRC II) in Computational Biology
2015-present Fellow, Canadian Institute for Advanced Research, Child and Brain Development

ACADEMIC AND TRAINING BACKGROUND

PDF 2014 Harvard Medical School (Dept. Immunology; mentor: Christophe Benoist)
PDF 2011-2014 Stanford University (Dept. Computer Science; mentor: Daphne Koller)
PhD 2006-2011 University of Toronto (Dept. Computer Science; advisor: Quaid Morris)
MS 2004-2006 Queen's University (Dept. Computing Sciences; advisor: Parvin Mousavi)
BSc 2001-2004 University of Toronto (Computer Science and Life Sciences)
1999-2001 Queen's University (Life Sciences) – transferred to UofT in 2001

SELECTED PUBLICATIONS

Trainees: underlined, co-first or co-senior authorship: *

Complete list: <https://scholar.google.ca/citations?user=nBL0J6kAAAAJ&hl=en>

Google scholar: H-index: 34; citations (all): 14470

(Selected) Under review:

Maslova, A., R. Ramirez, K. Ma, H. Schmutz, C. Wang, C. Fox, B. Ng, C. Benoist^{\$}, and **Mostafavi S^{\$}** (2019). Learning immune cell differentiation. (*available on bioRxiv*). (**Co-corresponding author**)

Patrick E, Taga M, Ergun A, Ng B, Casazza W, Cimpean M, Yung C, Schneider JA, Bennett DA, Gaiteri C, De Jager PL^{\$}, Bradshaw EM^{\$}, **Mostafavi S^{\$}**. (2019) Deconvolving the contributions of cell-type heterogeneity on cortical gene expression. (*available on bioRxiv*) (**Co-corresponding author**)

Graham E, Richmond PA, Tarailo-Graovac M, Engelke U, Kluijtmans L, Coene K, Wevers RA, Wasserman WW, van Karnebeek C, **Mostafavi S**. (2020). metPropagate: network-guided propagation of metabolomic information for prioritization of metabolic disease genes. (*minor revision; available on medRxiv*).

(Selected) Published:

1. Yoshida H, Lareau C, ..., **Mostafavi S***, Buenrostro J*, Benoist C*. The cis-Regulatory Atlas of the Mouse Immune System. (2019). *Cell*.
2. **Mostafavi S***, Gaiteri C*, Sullivan SE, ..., Young-Pearse TL, Bennett DA, De Jager PL. A molecular network of the aging human brain provides insights into the pathology and cognitive decline of Alzheimer's disease. (2018). *Nature Neuroscience*.

3. Ng B, White CC, Klein HU, Sieberts SK, McCabe C, Patrick E, Xu J, Yu L, Gaiteri C, Bennett DA, **Mostafavi S***, De Jager PL*(2017). An xQTL map integrates the genetic architecture of the human brain's transcriptome and epigenome. *Nature Neuroscience*.
4. **Mostafavi S***, Yoshida H*, Moodley D, ..., Mathis D*, Benoist C* and the Immunological Genome Project Consortium. (2016). Parsing the interferon transcriptional network and its disease associations. *Cell*.
5. Pierson E, the GTEx Consortium, Koller D, Battle A*, **Mostafavi S***. (2015) Sharing and specificity of co-expression networks across 35 human tissues. *PLOS Computational Biology*.
6. The GTEx Consortium. (2015) The Genotype-Tissue Expression (GTEx) pilot analysis: Multi-tissue gene regulation in humans. *Science*.
7. **Mostafavi S**, Ortiz-Lopez A, Hattori K, Pop C, Bogue M, Mathis D, Koller D, Benoist CB. (2014) Variation and genetic control of gene expression in primary immunocytes across inbred mouse strain. *Journal of Immunology*.
8. Raj T, Rothamel K, **Mostafavi S**, Ye C, Lee M, Replogle J, Von Korff A, Imboya S, McCabe C, Okada Y, Patsapolous N, Lee M, Wood I, Mathis D, Hafler D, Koller D, Regev A, Hacohen N, Benoist C*, Stranger BE*, De Jager PL*. (2014). Polarization of the effects of autoimmune and neurodegenerative risk alleles in leukocytes. *Science*.
9. Levinson DF, **Mostafavi S**, Milaneschi, Y, Rivera, M, Ripke S, Wray NR, Sullivan, PF. (2014) Genetics studies of major depressive disorder: Why are there no GWAS findings, and what can we do about it? *Biological Psychiatry*.
10. **Mostafavi S**, Battle A, Zhu X, ..., , Levinson D. (2013). Type I interferon signaling genes in recurrent major depression: increased expression detected by whole-blood RNA sequencing. *Molecular Psychiatry*
11. Battle A, **Mostafavi S**, Zhu X, Potash JB, Weissman MW, Mc-Cormick C, Haudenschild CD, Beckman K, Shi J, Mei R, Urban AE, Montgomery SB, Levinson D, Koller D. (2013). Characterizing the genetic basis of transcriptome diversity through RNA-sequencing of 922 individuals. *Genome Research*.
12. **Mostafavi S**, Battle A, Zhu X, Urban AE, Levinson D, Montgomery SB, Koller D. (2013). Normalizing RNA-sequencing data by modeling hidden covariates with prior knowledge. *PLOS ONE*.
13. Lonsdale J. et al., and the GTEx consortium. (2013). The Genotype-Tissue Expression (GTEx) project. *Nature Genetics*.
14. **Mostafavi S**, Goldenberg A, Morris Q. (2012). Labeling nodes using three degrees of propagation. *PLOS ONE*
15. **Mostafavi S**, Ray D, Warde-Farley D, Grouios C, Morris Q. (2008) GeneMANIA: A real-time multiple association network integration algorithm for predicting gene function. *Genome Biology*.

ACTIVE RESEARCH FUNDING

National Institutes of Health (NIH) U01 (sub-contract PI: Sara Mostafavi). *Multi-omic network-directed proteoforum discovery, dissection and functional validation to prioritize novel AD therapeutic targets*. 2018-2023.

NIH, R24 (sub-contract PI: Sara Mostafavi). *ImmGen: Gene expression in immune cells*. 2017-2022.

NIH, R01 (sub-contract PI: Sara Mostafavi). *Identifying the molecular systems networks, and key molecules that underlie cognitive resilience*. 2018-2022.

Canada Research Chair (Tier II) (PI). Chair in Computational Biology. 2015-2020.

Genome Canada, Large-Scale Applied Science Project (LSARP) (co-Investigator). *Childhood asthma and the microbiome—precision health for life: The Canadian Healthy Infant Longitudinal Development (CHILD) study*. 2018-2023.

Genome Canada, LSARP (co-Investigator). *Care4Rare-Solve: Multi-omic based diagnosis of rare disease*. 2018-2023.

Jacob Foundation, Catalyst Grant (co-PI). *Developing epigenetic measurements to forecast long-term benefits of early-life interventions*. 2018-2020.

Natural Sciences and Engineering Research of Canada (NSERC), Discovery Grant (PI). *Integrating multiple types of genomics data to identify meaningful associations*. 2016-2021.

Canadian Institute for Advanced Research (CIFAR) (PI). Child and Brain Development program. Research Fellow Funds. 2019-2024.

Canadian Institute for Advanced Research (CIFAR) (PI). Child and Brain Development program. *Cell type deconvolution algorithms*. 2015-2020.

Canadian Institute for Advanced Research (CIFAR) (PI). Catalyst Grant. *Construction of epigenetic growth chart*. 2016-2019.

Biogen Inc, SRA (PI). *Identifying cell-type specific eQTLs in the brain*. 2016-2019.

SELECTED INVITED PRESENTATIONS

- 2020 Department of Systems Biology, Harvard Medical School, Boston, MA.
- 2020 Department of Computer Science and Engineering, University of Washington, Seattle
- 2020 Department of Neurology, Columbia University, NY.
- 2019 Gladstone Institute, University of California, San Francisco (UCSF), CA.
- 2019 Nature Next Generation Genomics Conference at NYU, NY.
- 2019 Ludmir Center for Neuroinformatics, McGill University, Montreal.
- 2019 Center for Artificial Intelligence and Medicine, University of Montreal, Montreal.
- 2019 Department of Biostatistics, University of Washington, Seattle.
- 2019 Invited speaker, EMBL-EBI Industry and Academia Partnership: Neuroinformatics.
- 2019 Department of Computer Science, Boston University. Boston.
- 2018 Princess Margaret Hospital Research Center, Toronto, Canada
- 2018 Department of Human Genetics, University of California Los Angeles (UCLA), USA
- 2018 Women's Brain Initiative Symposium at Brigham Women's Hospital , Boston, USA
- 2018 CIFAR and Jacobs Foundation Meeting, Marbach Castle, Germany
- 2018 Invited speaker, Biogen Inc, Boston, MA, USA
- 2018 Statistical Seminar Series, McGill University, Montreal, Canada
- 2017 Invited speaker, Canadian Human & Statistical Genetics Meeting, Quebec City, Canada
- 2015 Mount Sinai School of Medicine, NY, USA
- 2015 Invited speaker, CIFAR meeting on Brain development, Vancouver, Canada
- 2015 Broad Institute, Population Genetics Meeting, Boston, MA

SELECTED PROFESSIONAL ACTIVITIES

Conference Chairing and Organization: Session Chair for Communities of Special Interest (COSI) at ISMB 2020; Track co-chair for ISMB 2019; Co-organizer for Machine Learning in Computational Biology (MLCB) workshop at NIPS 2013-2017, 2019; Co-organizer for Immunological Genome Project Computational Biology Meeting 2019; Co-organizer for BIRS

meeting on Statistical and Computational Challenges Arising from Ubiquitous Molecular Measurements 2020.

Conference Program Committee: ISMB 2013-2018; NIPS 2015-2019; PSB 2011-2016; RECOMB 2015-2019;

Grant Review Panels: Grant panel member for Canadian Institute for Health Research (CIHR) 2018, 2019; External reader for Natural Sciences and Engineering Research Council of Canada (NSERC) 2016, 2018, 2019; External reader for Research Council of UK 2017; Grant panel member for National Science Foundation (NSF) 2012; Ad-hoc reviewer for National Institute on Aging (NIA) 2017, 2018, 2019.

Journal Reviewing: Science, Nature Methods, Nature Communications, Genome Research, Genome Biology, Molecular Systems Biology, PLOS Genetics, PLOS Computational Biology, PLOS ONE, Annals of Applied Statistics, American Journal of Human Genetics, Biological Psychiatry, JAMA Neurology, Nucleic Acids Research, Bioinformatics, BMC Bioinformatics, Scientific Reports.

Education/Career Panels: Bioinformatics Training Program at UBC, Women in STEM 2018; Keynote for BC Children's Hospital Trainee Omics Group 2019; Medical Genetics Graduate Student Welcome Day, Career panel member 2019; Invited speaker for Invent The Future AI Scholar Program for high school girls 2019.