# **Hossein Sharifi-Noghabi**

Internship applicant/ Machine learning researcher/ Bioinformatics researcher Contact: (778) 885-4633/ Email: hsharifi@sfu.ca/ Website: https://hosseinshn.github.io/

### **Research interests**

Deep transfer learning, Unsupervised domain adaptation, adversarial transfer learning, few- and zero-shot learning, transfer learning in deep reinforcement learning, knowledge-based and interpretable models, applications in genomics and healthcare.

# **Education** Ph.D. in Computer Science (GPA: 4.08) Sep 2016-Present Simon Fraser University, Burnaby, BC, Canada. ☐ Supervisors: Prof. Martin Ester and Prof. Colin Collins Awards: Computing Science Graduate Fellowship (\$6.500, \$7.400, and \$8.000) ☐ Teaching Assistant: Introduction to computer programming I (Python) M.Sc. in Artificial Intelligence (GPA: 17.05/20) Sep 2012-Feb 2015 Ferdowsi University of Mashhad, Iran. ☐ Supervisor: Prof. H. Rajabi Mashhadi B.E. in Information Technology (GPA: 17.92/20) Sep 2008-July 2012 Sadjad University of Technology, Mashhad, Iran. ☐ Honor: Ranked 1st among 67 students of Information Technology **Selected publications** H. Sharifi-Noghabi, S. Peng, O. Zolotareva, C. Collins, M. Ester. (2019), "AITL: Adversarial Inductive Transfer Learning with input and output space adaptation for pharmacogenomics", MLCB 2019 and bioRxiv. Domain Knowledge-based Artificial Neural Network for drug response prediction", MLCB 2019.

- O. Snow, H. Sharifi-Noghabi, J. Lu, O. Zolotareva, M. Lee, M. Ester. (2019), "BDKANN Biological
- H. Sharifi-Noghabi, O. Zolotareva, C. C. Collins, M. Ester. (2019), "MOLI: Multi-Omics Late Integration with deep neural networks for drug response prediction", Bioinformatics 35 (14), i501-i509. Presented at ISMB/ECCB 2019. Received 1,500 USD travel award from the conference.
- H. Sharifi-Noghabi, et al.. (2019), "Deep Genomic Signature for early metastasis prediction in prostate cancer", Recomb-CCB 2019.
- M. Mohammadi, H. Sharifi-Noghabi, H. Rajabi Mashhadi, G. Hodtani. (2016), "Robust and stable gene selection via Maximum-Minimum Correntropy Criterion", Genomics (170) 83-87. (joint first authorship)

## W

Nork experience			
Research Assistant, Vancouver Prostate Centre (part time)		Mar 2017-Present	
	Obtained basic understanding of cancer biology-particularly properties of the project on metastasis prediction at 12th and Research Day		

	Participated in numerous research projects and grants as one	e of the bioinformaticians	
Rese	arch Assistant, Simon Fraser University (Full time)	Sep 2016-Present	
٥	Developed skills on deep unsupervised learning such as multi-task learning, and deep transfer learning (inductive and Designed and developed a novel methods in Pytorch and Kervia adversarial inductive transfer learning between cancer adapt the input domains in the genomic level, and 2) addifferent measures of drug response in cell lines and patients Finished two specializations on Coursera related to deel learning.	d adversarial transfer learning) as for drug response prediction cell lines and patients to: 1) apt the output spaces between.	
Resea	arch Intern, GenomeDx Inc. (Research collaboration)	Sep 2017-Jan 2018	
0	Developed a novel method in Tensorflow to predict metastasis in prostate cancer using Autoencoders and transfer learning on unlabelled and labelled genomic data and improved the accuracy compared to the state-of-the-art studies  Developed skills to analyze data via R such as survival analysis and data visualization  Obtained experience to collaborate with researchers with different backgrounds  Supervisors: Dr. Elai Davicioni, Nicholas Erho, and Dr. Yang (Seagle) Liu		
Select	ed course/personal projects during Ph.D.		
	natic chemical compounds design via Autoencoders using SMILES representation.  The goal was to generate new molecules via Variational Autoencoders. The hypothesis is that its learned representation is rich enough to sample and generate new molecules from it Used SMILES representations of existing chemical compounds and trained the Autoencoder on them		
Mini p	ersonal projects on Convolutional and Recurrent Neural N	letworks	
	Vision: Applied CNN to image verification and recognition tas Vision: Used neural style transfer to generate new arts NLP: Applied RNN models to synthesize Shakespeare's text NLP: Applied RNN to speech recognition and music synthesis		
Techr	nical and general skills		
Deep l Data s	mming languages: Python, C earning: Tensorflow, Pytorch, Keras cience: Matlab, R op and OS: Microsoft office, Latex, Photoshop, Windows, Ubunt	u	
Volun	teer positions		
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PC member—Machine learning in Computational Biology conference, Vancouver, Canada (2019). SFU Omics organizer—a group to provide an environment for students and other academics to come together to talk about their research related to Genomics, Proteomics, and Metabolomics. (2018-Present)