

Aryan Arbabi

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- EDUCATION
- ◇ **University of Toronto**, Toronto, Canada Sep. 2016 - present
PhD student in Computer Science
 - Also affiliated with Vector Institute, Sickkids hospital and University Health Network (UHN)
 - Thesis: “Machine Learning for Generating and Understanding Medical Text”
 - Adviser: Prof. Michael Brudno
 - ◇ **University of Toronto**, Toronto, Canada Sep. 2013 - Jun. 2015
MSc in Computer Science
 - Thesis: “Cell Free DNA Fragment-Size Distribution Analysis (FSDA) for Non-Invasive Prenatal CNV Prediction”
 - Adviser: Prof. Michael Brudno
 - ◇ **Sharif University of Technology**, Tehran, Iran Sep. 2009 - Aug. 2013
BSc in Computer Engineering
 - Thesis: “ARYANA: A Sequence Aligner for Next Generation Sequencing Data”
- WORK EXPERIENCE
- ◇ **Google Research**, Palo Alto, CA, USA July. 2019 - Oct. 2019
Research Intern
Research on generating descriptive text snippets conditioned on structured source data.
 - ◇ **Google Research**, New York, NY, USA Jun. 2018 - Sep. 2018
Software Engineering Intern
Research on Variational Autoencoders for unsupervised information extraction and representation learning for natural language.
 - ◇ **The Hospital for Sick Children**, Toronto, ON, Canada Jun. 2015 - Jun. 2020
Research Assistant
Research on Machine Learning and NLP methods to facilitate diagnosis of rare genetic diseases.
 - ◇ **University of Toronto, Department of Computer Science**, Toronto, ON, Canada Jun. 2015 - Aug. 2016
Research Software Engineer
Designed and developed Machine Learning methods for medical text processing.
- RESEARCH INTERESTS AND EXPERIENCE
- ◇ **Natural Language Processing.** Interested in techniques improving performance in NLU and NLG tasks with limited labeled data. Also interested in tasks involving graphs and natural language. Some of my projects in this direction:
 - Neural Concept Recognizer tool (NCR), improving detection of biomedical concepts mentioned in text, by utilizing the hierarchical relations between the concepts.
 - Unsupervised method based on Variational Autoencoders (VAEs), to detect informative words, such as proper nouns, in task oriented dialogue systems, without any annotated text.
 - BERT-based method that generates textual descriptions, conditioned on structured input such as knowledge graphs.
 - ◇ **Graph Neural Networks.** Other than tasks involving graphs and natural language, I have recently worked on utilizing Graph Neural Networks on protein-protein-interaction (PPI) networks, to predict functions for novel proteins. The task can be also viewed as a special case of Recommender Systems.
 - ◇ **Bayesian Deep Learning.** Interested in both theoretical and practical aspects of bayesian deep learning, approximate inference, deep generative models and information theoretic analysis of deep learning methods. Recently have been exploring using VAEs and EBMs for Recommender Systems.

- PUBLICATIONS
- ◇ **Aryan Arbabi** and Michael Brudno. Learning to propagate labels for protein function prediction. *under preparation*, 2021
 - ◇ **Aryan Arbabi**, Mingqiu Wang, Nan Du, Laurent El Shafey, and Izhak Shafran. R2d2: Relational text decoding with transformers. *arXiv preprint arXiv:2105.04645*, 2021
 - ◇ **Aryan Arbabi**, David R. Adams, Sanja Fidler, and Michael Brudno. Identifying clinical terms in free-text notes using ontology-guided machine learning. In *International Conference on Research in Computational Molecular Biology*, pages 19–34. Springer, 2019
 - ◇ Emily Fertig, **Aryan Arbabi**, and Alexander A Alemi. β -vae's can retain label information even at high compression. *Bayesian Deep Learning Workshop at NeurIPS*, 2018
 - ◇ Yanwei Xi[†], **Aryan Arbabi**[†], Amy JM McNaughton, Alison Hamilton, Danna Hull, Helene Perras, Tillie Chiu, Shawna Morrison, Claire Goldsmith, Emilie Creede, et al. Noninvasive prenatal detection of trisomy 21 by targeted semiconductor sequencing: A technical feasibility study. *Fetal Diagnosis and Therapy*, 2017
 - [†] These authors contributed equally to the work
 - ◇ **Aryan Arbabi**, Ladislav Rampásek, and Michael Brudno. Cell-free DNA fragment-size distribution analysis for non-invasive prenatal CNV prediction. *Bioinformatics*, 32(11):1662–1669, 2016
 - ◇ Ladislav Rampásek, **Aryan Arbabi**, and Michael Brudno. Probabilistic method for detecting copy number variation in a fetal genome using maternal plasma sequencing. *Bioinformatics*, 30(12):i212–i218, 2014
 - ◇ Milad Gholami[†], **Aryan Arbabi**[†], Ali Sharifi-Zarchi, Hamidreza Chitsaz, and Mehdi Sadeghi. ARYANA: Aligning Reads by Yet Another Approach. *BMC bioinformatics*, 15(Suppl 9):S12, 2014
 - [†] These authors contributed equally to the work
 - ◇ **Aryan Arbabi**, Milad Gholami, Mojtaba Varmazyar, and Shervin Daneshpajouh. Fast cpu-based DNA exact sequence aligner. In *Formal Methods and Models for Codesign (MEMOCODE), 2012 10th IEEE/ACM International Conference on*, pages 95–98. IEEE, 2012
- HONORS AND AWARDS
- ◇ **14th team** in the ACM-ICPC East Central North America Regional Contest, Nov. 2013, Windsor, Canada.
 - ◇ **1st team** in the Memocode 2012 Design Contest, Performance per Cost Section.
 - ◇ Recipient of the **grant for undergraduate studies** from the Iranian National Elites Foundation, for outstanding academic success, 2009 - 2013.
 - ◇ **Gold Medal** in the Iranian National Olympiad in Informatics (INOI), 2008.
- TEACHING EXPERIENCE
- ◇ **University of Toronto**, Toronto, Canada
Teaching Assistant Sep. 2013 - present
Courses: Probabilistic Learning and Reasoning, Introduction to Machine Learning, Natural Language Computing, Data Structures and Analysis, Introduction to Theory of Computation, Software Design, Introduction to Computer Programming
 - ◇ **Sharif University of Technology**, Tehran, Iran
Teaching Assistant Sep. 2011 - Apr. 2013
Courses: Design and Analysis of Algorithms, Theory of Languages and Automata, Discrete Structures, Advanced Programming
 - ◇ **Young Scholars Club**, Tehran, Iran
Teacher Summer 2010
Taught algorithms and programming to high school students accepted to Iranian Olympiad in Informatics (INOI) summer camp
- TECHNICAL SKILLS
- ◇ Python, TensorFlow, PyTorch, Java, C/C++