

INTERPRETABLE DEEP LEARNING FOR CANCER RESEARCH

SEPTEMBER 20-21, 2018

NATIONAL INSTITUTES OF HEALTH BUILDING 31 C-WING 6TH FLOOR, ROOM 6 31 CENTER DRIVE BETHESDA, MARYLAND 20892

INTRODUCTION OF THE WORKSHOP

Recent developments in convolutional neural networks and related machine learning techniques commonly referred to as deep learning (DL) have made breakthroughs in computer vision, robotic control, machine translation, voice recognition, gaming, and many other areas. While DL has seen rapid adoption by various cancer imaging communities, its use by the broader cancer biology community has been less marked. This can be attributed, at least in part, to a limitation frequently referred to as the "black box", where DL models can make correct predictions but without association to underlying mechanisms for biological interpretation. This workshop will bring together DL researchers, cancer systems biologists, and computational biologists to discuss challenges and opportunities to develop interpretable DL methods that can be applied to cancer biology investigations for knowledge generation. The workshop aims to enumerate significant hurdles and identify potential pathways to promote innovations in DL method development, with the goal of overcoming existing limitations and making the DL approach most impactful in cancer research.

DISCUSSION TOPICS OF WORKSHOP SESSIONS:

Session 1. **Transferability of** *tabula rasa* **learning methods to biomedical problems.** Most deep learning models developed in the biomedical domain rely on benchmarked datasets for supervised training. Entirely rule-based learning without human knowledge or data has demonstrated success in competing with traditional supervised learning models in certain areas such as gaming. This session will focus on discussing the transferability of deep learning algorithms that learn *tabula rasa* to solve biological problems.

Session 2. **Transferability of biomedical domain knowledge to deep learning methods development**. The original design of deep learning neural networks was inspired by the structure of the human brain. With the ever-increasing sophistication of deep learning network development, biological knowledge can contribute to the design of new topology of neural networks, new training methods, and new ways of integration with other biologically informed machine learning algorithms. This session focuses on discussing avenues to integrate biological knowledge in designing new deep learning methods.

Session 3. Challenges and solutions associated with noisy, heterogeneous, and limited amount of data. Biological data are intrinsically noisy, heterogeneous, and are often distributed and insulated with limited access. This session will focus on discussing strategies to improve model quality with limited amounts of training data and/or data quality considerations.

Session 4. **Interpretability of deep learning models**. This session will focus on discussing strategies to open up the "black box" and make output interpretable for mechanistic insight and knowledge generation.

Session 5. **Model validation, sharing, and comparison**. This session will focus on discussing strategies and methods for model validation, model sharing, and model comparison so that we can learn from each other and improve as a field. Key resource needs will also be discussed at this session.

ORGANIZING COMMITTEE:

Trey Ideker, Ph.D.	University of California, San Diego
Claire Tomlin, Ph.D.	University of California, Berkeley
Jennifer Couch, Ph.D.	National Cancer Institute
Jerry Li, M.D., Ph.D.	National Cancer Institute
David Miller, Ph.D.	National Cancer Institute

AGENDA: THURSDAY, SEPTEMBER 20, 2018

8:15 am	Shuttle departs hotel for NIH	
9:00 am – 9:15 am	WELCOME AND INTRODUCTION OF THE WORKSHOP	
	Dinah Singer, Ph.D.	
	Acting Deputy Director, National Cancer Institute	
	Trey Ideker, Ph.D.	
	Professor of Medicine, University of California, San Diego	
	Claire Tomlin, Ph.D.	
	Professor of Electrical Engineering and Computer Sciences, University of	
	California, Berkeley	
9:15 am – 10:45 am	OPENING KEYNOTE PRESENTATIONS	
9:15 am – 9:55 am	Overview of new deep learning methods	
	Rick Stevens, Ph.D.	
	Argonne National Laboratory	
9:55 am – 10:30 am	Interpretable deep learning for biological insights	
	Trey Ideker, Ph.D.	
	Professor of Medicine, University of California, San Diego	
10:30 am – 10:45 am	NCI Director's remarks	
	Norman Sharpless, M.D.	
	Director, National Cancer Institute	
10:45 am – 11:00 am	BREAK	

11:00 am – 12:40 pm	SESSION 1: TRANSFERABILITY OF <i>TABULA RASA</i> LEARNING METHODS TO BIOMEDICAL PROBLEMS		
	Panel members: Rick Stevens, Ph.D. (Chair) Ziv Bar-Joseph, Ph.D. Anna Goldenberg, Ph.D. Shun Miao, Ph.D.	Argonne National Laboratory Carnegie Mellon University University of Toronto NVidia	
11:00 am – 11:50 am 11:50 am – 12:40 pm	Short presentations from par Panel discussion	nel members	
12:40 pm – 1:40 pm	LUNCH (ON OWN)		
1:40 pm – 3:30 pm	SESSION 2: TRANSFERABILITY OF BIOMEDICAL DOMAIN KNOWLEDGE TO DEEP LEARNING METHODS DEVELOPMENT		
	Panel members: Chris Sander, Ph.D. (Chair) Anthony Gitter, Ph.D. Jinbo Xu, Ph.D. Alexander Anderson, Ph.D.	Dana-Farber Cancer Institute University of Wisconsin Madison Toyota Technological Institute at Chicago Moffitt Cancer Center	
1:40 pm – 2:40 pm 2:40 pm – 3:30 pm	Short presentations from panel members Panel discussion		
3:30 pm – 3:45 pm	BREAK		
3:45 pm – 5:35 pm	SESSION 3: CHALLENGES AND SOLUTIONS ASSOCIATED WITH NOISY, HETEROGENEOUS, AND LIMITED AMOUNT OF DATA		
3:45 pm – 4:45 pm 4:45 pm – 5:35 pm	Panel members: Claire Tomlin, Ph.D. (Chair) Anshul Kundaje, Ph.D. Young Hwan Chang, Ph.D. Kristin Swanson, Ph.D. Short presentations from pan Panel discussion	University of California, Berkeley Stanford University Oregon Health & Science University Mayo Clinic nel members	
E-25 nm			
5:45 pm	Shuttle from NIH back to hotel		
7:00 pm – 9:00 pm	Group dinner at Chef Tony's restaurant 4926 St Elmo Avenue, Bethesda, Md		

AGENDA: FRIDAY SEPTEMBER 21, 2018

8:15 am	Shuttle departs hotel for NIH		
9:00 am – 10:50 am	SESSION 4: INTERPRETABILITY OF DEEP LEARNING MODELS		
	Panel members: Regina Barzilay, Ph.D. (Chair) Elias Bareinboim, Ph.D. Xinghua Lu, M.D., Ph.D. Trey Ideker, Ph.D.	Massachusetts Institute of Technology Purdue University University of Pittsburgh University of California, San Diego	
9:00 am – 10:00 am 10:00 am – 10:50 am	Short presentations from panel members Panel discussion		
10:50 am – 11:05 am	BREAK		
11:05 am – 12:50 pm	SESSION 5: MODEL VALIDATIO	N. SHARING. AND COMPARISON	
11:05 am – 12:05 pm 12:05 pm– 12:50 pm 12:50 pm– 1:50 pm	Panel members: Peter Sorger, Ph.D. (Chair) Emek Demir, Ph.D. Doron Levy, Ph.D. Ronald Summers, M.D., Ph.D. Short presentations from panel Panel discussion	Harvard Medical School Oregon Health & Science University University of Maryland College Park NIH Clinical Center	
1:50 pm – 3:00 pm	Chairs: Trey Ideker and Claire Tomlin		
1:50 pm – 2:40 pm	Session summaries Session Chairs		
2:40 pm – 3:00 pm	Summary of the workshop and closing remarks Trey Ideker and Claire Tomlin		
3:00 pm	ADJOURN		