

# 3rd Course on Computational Systems Biology of Cancer: Promises of Artificial Intelligence

## September 28 - October 2, 2020

VIRTUAL

EMBO / FEBS / Institut Curie Training Unit International Course

VIRTUAL

Monday, September 28 <sup>th</sup>		Session 1: Machine learning in prior knowledge applications for data analysis
Chairs: TBA		
08:50 09:00	<b>Emmanuel Barillot</b> Institut Curie, FR	Welcome and opening remarks
09:00 10:00	<b>Andrei Zinovyev</b> Institut Curie, FR	Didactic introductory lecture on AI approaches in systems biology
10:00 10:30	Coffee break / Meet a speaker in a virtual discussion room	
10:30 11:30	Selected Talks from Abstracts	15 minutes/presentation
11:30 12:30	Flash presentations of projects by post-docs	7 minutes/presentation
12:30 13:30	Lunch / See posters online	
Monday, September 28 <sup>th</sup>		Session 2: Network biology in Artificial Intelligence era
Chairs: TBA		
13:30 14:30	<b>Laura Cantini</b> IBENS-ENS, FR	Benchmarking of computational approaches to multi-omics data analysis
14:30 15:30	<b>Nataša Pržulj</b> Barcelona Supercomputing Center, ES	Analyzing Network Data in Biology and Medicine
14:30 15:30	Flash presentations of projects by PhD students	5 minutes/presentation
17:30 19:00	Welcome cocktail / Virtual Apéro	

Tuesday, September 29 <sup>th</sup>		Session 3: Patient stratification and disease classification using Artificial Intelligence methods (1)
Chairs: TBA		
09:00 10:00	<b>Valentina Boeva</b> ETH, Zurich, CH	Machine learning methods for the detection of intratumoral heterogeneity in cancer
10:00 10:30	Coffee break / Meet a speaker in a virtual discussion room	
10:30 11:30	<b>David Wedge</b> The Big Data Institute, UK	Stratification of prostate cancers through the application of machine learning to sequencing data
11:30 12:30	Selected Talks from Abstracts	15 minutes/presentation
12:30 13:30	Lunch / See posters online	
Tuesday, September 29 <sup>th</sup>		Session 4: Patient stratification and disease classification using Artificial Intelligence methods (2)
Chairs: TBA		
13:30 14:30	<b>Susana Vinga</b> Instituto Superior Técnico, PT	Cancer classification and survival analysis through sparse optimization and network-based regularisers
14:30 15:00	Coffee break / Meet a speaker in a virtual discussion room	
15:00 16:00	<b>Nataliya Sokolovska</b> University Pierre et Marie Curie, FR	Interpretable models for clinical scores
16:00 17:30	Flash presentations of projects by post-docs	7 minutes/presentation
17:30 18:00	COVID19 and cancer / Virtual round table discussion	

Wednesday, September 30 <sup>th</sup>		Session 5: Machine Learning applications in precision medicine (1)
Chairs: TBA		
09:00 10:00	<b>Chloé-Agathe Azencott</b> Centre for Computational Biology, MINES ParisTech, FR	TBA
10:00 10:30	Coffee break / Meet a speaker in a virtual discussion room	
10:30 11:30	<b>Yves Moreau</b> KULeuven, BE	Bayesian matrix factorization and deep learning for drug discovery and precision medicine
11:00 12:30	Flash presentations of projects by PhD students	5 minutes/presentation
12:30 13:30	Lunch / See posters online	
Wednesday, September 30 <sup>th</sup>		Session 6: Machine Learning applications in precision medicine (2)
Chairs: TBA		
13:30 14:30	<b>Jean-Philippe Vert</b> Google, FR	Robust machine learning for cancer precision medicine
14:30 15:00	<b>Olli Kallioniemi</b> SciLifeLab, SE	AI approaches in Precision Cancer Medicine
15:00 16:00	Coffee break / Meet a speaker in a virtual discussion room	
16:00 17:00	<b>Lodewyk Wessels</b> Netherlands Cancer Institute, NL	Mapping cellular networks in cancer
17:00 18:00	Selected Talks from Abstracts	15 minutes/presentation

<b>Thursday, October 1st</b>		<b>Session 7: Tumor genetics and epigenetics (1)</b>
Chairs: TBA		
09:00 10:00	<b>Christina Leslie</b> Memorial Sloan Kettering Cancer Center, USA	<b>Machine learning in regulatory genomics and systems biology</b>
10:00 10:30	Coffee break / Meet a speaker in a virtual discussion room	
10:30 11:30	<b>Andrew Teschendorff</b> University College, UK	<b>Machine Learning approaches for large scale multi-dimensional cancer omics data analysis</b>
11:30 12:30	<b>Julio Saez-Rodriguez</b> Heidelberg University, DE	<b>Supporting Machine Learning with Prior Biological Knowledge to extract insight from omics data</b>
12:30 13:30	Lunch / See posters online	
<b>Thursday, October 1st</b>		<b>Session 8: Tumor genetics and epigenetics (2)</b>
Chairs: TBA		
13:30 14:30	<b>Touati Benoukraf</b> MUN, CA	<b>DNA methylation and Cancer</b>
14:30 15:30	<b>List of speakers TBA</b>	<b>Career development WS: Invited speakers, pharma and scientific editing representatives, platform managers speaking on their career</b>
15:30 16:00	Coffee break / Meet a speaker in a virtual discussion room	
16:00 16:30	Master student's journal club	Presenentation of 4-5 milestone papers
16:30 17:30	Tutorial/Demo	TBA

<b>Friday, October 2nd</b>		<b>Session 9: Bioimage Informatics in cancer research (1)</b>
Chairs: TBA		
09:00 10:00	<b>Nasir Rajpoot</b> University of Warwick, UK	<b>The Promise of Computational Pathology</b>
10:00 10:30	Coffee break / Meet a speaker in a virtual discussion room	
10:30 11:30	<b>Florian Markowitz</b> University of Cambridge, UK	<b>Image analysis of the tumour tissue and early detection by AI</b>
11:30 12:30	Master student's journal club	Presenentation of 4-5 milestone papers
12:30 13:30	Lunch / See posters online	
<b>Friday, October 2nd</b>		<b>Session 10: Bioimage Informatics in cancer research (2)</b>
Chairs: TBA		
13:30 14:30	<b>Thomas Walter</b> Centre for Computational Biology, MINES ParisTech, FR	<b>Exploring the spatial aspects of gene expression</b>
14:30 15:30	<b>Emma Lundberg</b> SciLifeLab, SE	<b>AI for improved spatial proteomics</b>
15:30 16:00	Coffee break / Meet a speaker in a virtual discussion room	
16:00 17:30	Master student's journal club	Presenentation of 4-5 milestone papers
17:30 18:00	Closing remarks, prizes for presentations and posters	
17:30 19:00	Farewell cocktail / Virtual Apéro	