



COCHAIRS:

Valentina Boeva, ETH Zurich, Zurich, Switzerland

Marzyeh Ghassemi, Massachusetts Institute of Technology, Cambridge, Massachusetts

Benjamin Haibe-Kains, UHN Princess Margaret Cancer Centre, Toronto, Ontario, Canada

Bo Wang, University of Toronto and Vector Institute, Toronto, Ontario, Canada

THURSDAY, JULY 10

Registration

4-8:30 P.M. | Grand Salon Opera Foyer

Welcome and Keynote Lecture

6:30-7:15 P.M. | Grand Salon Opera AB

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| 6:30-6:35 | Welcome and Introduction of Keynote Speaker
Valentina Boeva, ETH Zurich, Zurich, Switzerland |
| 6:35-7:05 | Computational approaches to mapping cells, tissues and tumor progression, where does AI help?
Dana Pe'er, Memorial Sloan Kettering Cancer Center, New York, New York |
| 7:05-7:15 | Discussion / Q&A |

Poster Session A / Opening Reception

7:30-9 P.M. | Soprano ABC and Foyer

FRIDAY, JULY 11

Continental Breakfast and Networking Roundtables

7-8 A.M. | Grand Salon Foyer and Grand Salon Opera C

Keynote Lecture

8-8:45 P.M. | Grand Salon Opera AB

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| 8:00-8:05 | Introduction of Keynote Speaker
Benjamin Haibe-Kains, UHN Princess Margaret Cancer Centre, Toronto, Ontario, Canada |
| 8:05-8:35 | Empowering biomedical discovery with "AI scientists"
Marinka Zitnik, Harvard University, Boston, Massachusetts |
| 8:35-8:45 | Discussion / Q&A |

American Association for Cancer Research® Special Conference in Cancer Research

Artificial Intelligence and Machine Learning

July 10-12, 2025 | Montreal, CA



Plenary Session 1: ML/AI for Drug Discovery, Repurposing, and Response Prediction

8:50-10:10 A.M. | Grand Salon Opera AB

Session Chair: Benjamin Haibe-Kains, UHN Princess Margaret Cancer Centre, Toronto, ON, Canada

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| 8:50-9:15 | Supporting anticancer drug discovery by knowledge graph mining and cancer target-focused cheminformatics modeling
Alexander Tropsha, University of North Carolina, Chapel Hill, North Carolina |
| 9:15-9:20 | Discussion / Q&A |
| 9:20-9:45 | Virtual cells and digital twins - AI in personalized oncology
Charlotte Bunne, Swiss Federal Institute of Technology Lausanne (EPFL), Lausanne, Switzerland |
| 9:45-9:50 | Discussion / Q&A |
| 9:50 –10:05 | AI-Predict: Artificial intelligence-mediated drug synergy prediction and validation in cancer models*
Alicia Pliego, University of Zurich, Zurich, Switzerland |
| 10:05-10:10 | Discussion / Q&A |

Coffee Break

10:10-10:25 A.M. | Grand Salon Foyer

Plenary Session 2: ML/AI for Genomic and Temporal Predictions

10:25-11:45 A.M. | Grand Salon Opera AB

Session Chair: Bo Wang, University of Toronto and Vector Institute, Toronto, Ontario, Canada

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| 10:25-10:50 | Spatial cell-state dynamics in the tumor microenvironment
Samantha Riesenfeld, University of Chicago, Pritzker School of Molecular Engineering, Chicago, Illinois |
| 10:50-10:55 | Discussion / Q&A |
| 10:55-11:20 | Illuminating the Dark Cancer Genome using Foundation Models
Raul Rabadan, Columbia University Irving Medical Center, New York, New York |
| 11:20-11:25 | Discussion / Q&A |
| 11:25-11:40 | Genomic alterations associate with distinct tumor dynamics in NSCLC: A genomics-informed Bayesian hierarchical modeling approach*
Nikolaos Dimitriou, Memorial Sloan Kettering Cancer Center, New York, New York |
| 11:40-11:45 | Discussion / Q&A |

Break

11:45 A.M.-12 P.M. | Grand Salon Foyer

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*-short talk from proffered abstracts

AI Synergy Forum: Fostering Collaboration Through Innovation

12-12:45 P.M. | Grand Salon Opera AB

Session Chair: Benjamin Haibe-Kains, UHN Princess Margaret Cancer Centre, Toronto, ON, Canada

12-12:05	Theme 1: Open science: Data and code sharing + reusability Shirin A. Enger, McGill University, Montreal, Quebec, Canada
12:05-12:10	Theme 2: AI using non/minimally invasive modalities for cancer applications Florian Markowetz, Cancer Research UK Cambridge Institute, Cambridge, UK
12:10-12:15	Theme 3: Monitoring model performance/drift in research and clinical settings Elana J. Fertig, University of Maryland, Baltimore, Maryland
12:15-12:20	Theme 4: The case for collaboration for real world impact Skye Bork, PACT AI
12:20-12:45	Panel Discussion

Panelists:

Skye Bork, PACT AI

Jian Carrot-Zhang, Memorial Sloan Kettering Cancer Center, New York, New York

Shirin A. Enger, McGill University, Montreal, Quebec, Canada

Elana J. Fertig, University of Maryland, Baltimore, Maryland

Florian Markowetz, Cancer Research UK Cambridge Institute, Cambridge, UK

Raul Rabadan, Columbia University Irving Medical Center, New York, New York

Free time / Lunch on Own

12:45-2:15 P.M.

Plenary Session 3: Multimodal AI for Clinical Applications in Oncology

2:15-3:35 P.M. | Grand Salon Opera AB

Session Chair: Dana Pe'er, Memorial Sloan Kettering Cancer Center, New York, New York

2:15-2:40	Generative AI for improved digital pathology model predictions and interpretability [R] Christina Curtis, Stanford University, Stanford, California, USA
2:40-2:45	Discussion / Q&A
2:45-3:10	Alignment and integration of spatial multi-omics tumor profiles Ben Raphael, Princeton University, Princeton, New Jersey
3:10-3:15	Discussion / Q&A
3:15-3:30	Inferring genomic properties and histologic subtypes of solid tumors from H&E whole-slide images*

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3:30-3:35 Kevin Boehm, MSKCC, New York, New York
 Discussion / Q&A

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 [R] – Remote Presentation

Break

3:35-3:50 P.M. | Grand Salon Foyer

Plenary Session 4: Bias and Fairness in AI Research and Deployed Models

3:50-5:10 P.M. | Grand Salon Opera AB

Session Chair: Benjamin Haibe-Kains, UHN Princess Margaret Cancer Centre, Toronto, Ontario, Canada

3:50-4:15	Uncovering and mitigating bias in auto-segmentation models in radiation oncology Benjamin Haibe-Kains, UHN Princess Margaret Cancer Centre, Toronto, Ontario, Canada
4:15-4:20	Discussion / Q&A
4:20-4:45	Beyond assistance: Rethinking AI-human integration in radiology [R] Pranav Rajpurkar, Harvard University, Boston, Massachusetts
4:45-4:50	Discussion / Q&A
4:50-5:05	A practical framework for operationalizing responsible and equitable AI in healthcare: Tackling bias, inequity, and implementation challenges* Benjamin Grant, Cancer Digital Intelligence, Princess Margaret Cancer Centre, Toronto, Ontario, Canada
5:05-5:10	Discussion / Q&A

Break

5:10-5:25 P.M. | Grand Salon Foyer

Plenary Session 5: Cancer Applications of Foundation Models

5:25-6:45 P.M. | Grand Salon Opera AB

Session Chair: Valentina Boeva, ETH Zurich, Zurich, Switzerland

5:25-5:50	Cancer-specific foundation models: Friend or foe in healthcare AI? Valentina Boeva, ETH Zurich, Zurich, Switzerland
5:50-5:55	Discussion / Q&A
5:55-6:20	Evaluating and overseeing large language models for oncology Danielle S. Bitterman, Harvard Medical School, Boston, Massachusetts
6:20-6:25	Discussion / Q&A



- 6:25-6:40 Learning the language of somatic mutations: A large language model approach to precision oncology*
John-William Sidhom, Weill Cornell Medicine, New York, New York
- 6:40-6:45 Discussion / Q&A

*-short talk from proffered abstracts
[R] – Remote Presentation

Poster Session / Reception
7-9 P.M. | Soprano ABC and Foyer

SATURDAY, JULY 12

Continental Breakfast and Networking Roundtables
7-8 A.M. | Grand Salon Foyer and Grand Salon Opera C

Keynote Lecture 3: AI for Image Analysis
8-8:45 P.M. | Grand Salon Opera AB

- 8-8:05 Introduction of Keynote Speaker
Bo Wang, University of Toronto and Vector Institute, Toronto, Ontario, Canada
- 8:05-8:35 Accelerating oncology drug discovery with the power of microscopy and AI
Anne Carpenter, Broad Institute, Cambridge, Massachusetts
- 8:35-8:45 Discussion / Q&A

Plenary Session 6: ML/AI for Single Cell and Spatial Data, H&E, and Molecular Biomarker Discovery
8:50-10:10 A.M. | Grand Salon Opera AB
Session Chair: Ben Raphael, Princeton University, Princeton, NJ

- 8:50-9:15 **Forecasting tumor-immune dynamics in pancreatic cancer**
Elana J. Fertig, University of Maryland, Baltimore, Maryland
- 9:15-9:20 Discussion / Q&A
- 9:20-9:45 Towards fast and accessible precision oncology
Ruppin Eytan, National Cancer Institute, Bethesda, Maryland
- 9:45-9:50 Discussion / Q&A
- 9:50-9:57 Analysis of pathologists' intraobserver, interobserver and AI agreement in breast cancer HER2 scoring: AI-assessed intra-sample tumor heterogeneity relates to lower agreement among pathologists and with AI*
Pedro Simonis S. M. Ferrari, D'Or Institute of Research & Education, São Paulo, Brazil
- 9:57-10 Discussion / Q&A



- 10-10:07 Path2Space: An AI approach for cancer biomarker discovery via histopathology inferred spatial transcriptomics*
Emma Campagnolo, National Cancer Institute (NIH), Bethesda, Maryland, United States
- 10:07-10:10 Discussion / Q&A
- 10:10-10:20 Automated segmentation pipeline for radiological imaging using UniverSeg and similarity-guided support set retrieval*
Niket Patel, Drexel University College of Medicine, Philadelphia, Pennsylvania
- 10:20-10:25 Discussion / Q&A

*-short talk from proffered abstracts

Coffee Break

10:25-10:45 A.M. | Grand Salon Foyer

Plenary Session 7: ML/AI for Radiological Imaging

10:45 A.M.-11:50 A.M. | Grand Salon Opera AB

Session Chair: Anne Carpenter, Broad Institute, Cambridge, Massachusetts

- 10:45-11:10 **ML/AI for radiological imaging**
Catherine Coolens, University Health Network, Toronto, Ontario, Canada
- 11:10-11:15 Discussion / Q&A
- 11:15-11:40 **Integration of pathology with radiological imaging to extract biological signals for treatment guidance**
Caroline Chung, UT MD Anderson Cancer Center, Houston, Texas
- 11:40-11:45 Discussion / Q&A

Plenary Session 8: How do we make AI research better for society!

11:50-1:20 P.M. | Grand Salon Opera AB

Session Chair: Elana J. Fertig, University of Maryland, Baltimore, Maryland

- 11:50-12:15 **Research in ethical machine learning and health [R]**
Marzyeh Ghassemi, Massachusetts Institute of Technology, Cambridge, Massachusetts
- 12:15-12:20 Discussion / Q&A
- 12:20-12:45 **Building, scaling and using AI engines**
Jeff Leek, Fred Hutchinson Cancer Center, Seattle, Washington
- 12:45-12:50 Discussion / Q&A
- 12:50-1:15 The impact of bringing complex data to the point of care
Casey Greene, University of Colorado, Aurora, Colorado
- 1:15-1:20 Discussion / Q&A

Closing Remarks and Departure

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1:20 P.M.

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