

AACR Virtual Meeting Advances in Malignant Lymphoma August 17-19, 2020



AACR Virtual Meeting

Advances in Malignant Lymphoma: Maximizing the Basic-Translational Interface for Clinical Application

In cooperation with the International Conference on Malignant Lymphoma (ICML)

August 17–19, 2020 | Virtual Meeting

Monday, August 17

Welcome and Opening Keynote Address

10:30-11:15 a.m. Channel 1

10:30–10:35 a.m. Welcome and Keynote Introduction

Ari M. Melnick, Weill Cornell Medical College, New York, NY

10:35–11 a.m. Rogue clones of self-reactive B cells with lymphoma driver mutations:

A view of lymphoma precursors

Christopher Goodnow, Garvan Institute of Medical Research, Darlinghurst,

NSW. Australia

11–11:15 a.m. Discussion / Q&A

Break

11:15-11:30 a.m.

Session 1: Teaching T-Cells New Tricks: Why Immunotherapy Fails in Lymphoma and How Do We Fix It?

Moderator: Helen E. Heslop, Baylor College of Medicine, Houston, TX 11:30 a.m.–1:05 p.m.

Channel 1

11:30–11:35 a.m. Moderator Introduction

11:35–11:50 a.m. The bad guys of follicular lymphoma microenvironment: Putative

impact on therapy failure?

Karin Tarte, Université de Rennes, Rennes, France

11:50 a.m.-12:05 p.m. Immune exhaustion and suppression in lymphoma

Stephen M. Ansell, Mayo Clinic College of Medicine, Rochester, MN

12:05–12:20 p.m. Optimizing immunostimulatory antibodies for cancer immunotherapy

Mark Cragg, University of Southampton, Southampton, United Kingdom

12:20–12:35 p.m. The immune cell microenvironment of classical Hodgkin lymphoma by

single-cell analysis

Christian Steidl, BC Cancer, Vancouver, BC, Canada

12:35–1:05 p.m. Discussion / Q&A

Break

1:05-1:20 p.m.

Session 2: WHO vs. How? Changing Paradigms in Lymphoma Classification

Moderator: Georg Lenz, University of Münster, Münster, Germany 1:20–2:55 p.m.

Channel 1

1:20–1:25 p.m. Moderator Introduction

1:25–1:40 p.m. **Molecular classification of T-cell lymphomas**

Keisuke Kataoka, National Cancer Center, Japan, Tokyo, Japan

1:40–1:55 p.m. The role of gene expression in the classification of aggressive B-cell

lymphoma

David W. Scott, BC Cancer, Vancouver, BC, Canada

1:55–2:10 p.m. Molecular classification of diffuse large B-cell lymphoma

Bjoern Chapuy, University of Göttingen, Göttingen, Germany

2:10–2:25 p.m. Microenvironmental signatures reveal biological and clinical subtypes

of diffuse large B-cell lymphoma

Leandro C. Cerchietti, Weill Cornell Medical College, New York, NY

2:25–2:55 p.m. Discussion / Q&A

Break

2:55-3:10 p.m.

Session 3: Towards a Better Understanding of Adolescent Young Adult Lymphoma

Moderator: Sonali M. Smith, University of Chicago, Chicago, IL 3:10–4:20 p.m.

Channel 1

3:10–3:15 p.m. Moderator Introduction

3:15–3:30 p.m.	AYA lymphomas: Bridging the divide Kara Kelly, Roswell Park Comprehensive Cancer Center, Buffalo, NY
3:30–3:45 p.m.	Genomics of common AYA lymphomas: Hodgkin lymphoma and primary mediastinal B-cell lymphoma Lisa G. Giulino-Roth, Weill Cornell Medical College, New York, NY
3:45–4:00 p.m.	Unique features of pediatric-type follicular lymphoma Abner Louissaint, Massachusetts General Hospital, Boston, MA
4:00–4:20 p.m.	Discussion / Q&A

Session 4: How Aberrant Metabolism Fuels Lymphoma

Moderator: Francesco Bertoni, Institute of Oncology Research, Bellinzona, Switzerland 3:10-4:05 p.m. Channel 2

3:10–3:15 p.m.	Moderator Introduction
3:15–3:30 p.m.	Oncogenic Rag GTPase signaling links cellular nutrients with FL microenvironment Alejo Efeyan, Spanish National Cancer Research Center, CNIO, Madrid, Spain
3:30–3:45 p.m.	Metabolic control of lymphomagenesis Andrew Michael Intlekofer, Memorial Sloan Kettering Cancer Center, New York, NY
3:45-4:05 p.m.	Discussion / Q&A

Tuesday, August 18

Session 5: New CARs and Fine-Tuning Patient Selection for Therapy
Moderator: Renier J. Brentjens, Memorial Sloan Kettering Cancer Center, New York, NY 10-11:20 a.m. Channel 1

10–10:05 a.m.	Moderator Introduction
10:05–10:20 a.m.	CARs and armored CARs: Improving CAR T cell therapy for cancer Renier J. Brentjens
10:20–10:35 a.m.	CAR T-cells for T-cell lymphoma Maksim Mamonkin, Baylor College of Medicine, Houston, TX

10:35–10:50 a.m. **Determinants of CAR-T outcomes: Host, tumor, and product**

Sattva S. Neelapu, The University of Texas MD Anderson Cancer Center,

Houston, TX

10:50-11:20 a.m. Discussion / Q&A

Break

11:20-11:50 a.m.

Session 6: Biology-Based Clinical Trials: A Myth or Hope for Reality?

Moderator: John F. Seymour, Peter MacCallum Cancer Centre, Melbourne, VIC, Australia 11:50 a.m.–1:25 p.m.

Channel 1

11:50–11:55 a.m. Moderator Introduction

11:55 a.m.-12:10 p.m. Can we hope for success in biologically stratified DLBCL studies?

Andrew John Davies, University of Southampton School of Medicine,

Southampton, United Kingdom

12:10–12:25 p.m. Why good trials fail: Impact of biomarker requirement on the selection

bias of study patients

Grzegorz S. Nowakowski, Mayo Clinic College of Medicine and Science,

Rochester, MN

12:25–12:40 p.m. Biomarker informed studies in peripheral T-cell lymphoma

Steven M. Horwitz, Memorial Sloan Kettering Cancer Center, New York, NY

12:40–12:55 p.m. **Precision treatment of lymphoma using genomics**

Louis M. Staudt, National Cancer Institute, Bethesda, MD

12:55–1:25 p.m. Discussion / Q&A

Break

1:25-1:40 p.m.

Session 7: Hacking the Lymphoma Epigenome to Create Better Therapies

Moderator: Ari M. Melnick, Weill Cornell Medical College, New York, NY 1:40–3:15 p.m.

Channel 1

1:40–1:45 p.m. Moderator Introduction

1:45–2:00 p.m. The role of EZH2 gain-of-function mutations in lymphoma

Wendy Béguelin, Weill Cornell Medical College, New York, NY

2:00–2:15 p.m. Cooperative epigenetic interactions in normal and transformed germinal center B cells
 Laura Pasqualucci, New York Presbyterian Hospital Herbert Irving Comprehensive Cancer Center, New York, NY

 2:15–2:30 p.m. CREBBP: Not all mutations are created equal Michael R. Green, The University of Texas MD Anderson Cancer Center, Houston, TX
 2:30–2:45 p.m. What do the results from the trials tell us?
 Peter W. M. Johnson, University of Southampton, Southampton, United Kingdom

 2:45–3:15 p.m. Discussion / Q&A

Break

3:15-3:30 p.m.

Session 8: Cancer Health Disparities in Lymphoma: How Do We Improve Outcomes?

Moderator: Christopher R. Flowers, The University of Texas MD Anderson Cancer Center, Houston, TX 3:30–4:40 p.m.

Channel 1

	3:30–3:35 p.m.	Moderator Introduction
survivors of lymphoma Theresa H. M. Keegan, University of California Davis, Sacramento, CA 4:05–4:20 p.m. Genome-estimated African ancestry is associated with distinct tumor mutations and poorer survival in patients with diffuse large B-cell lymphoma Christopher R. Flowers	3:35–3:50 p.m.	Risk disparities in lymphoid cancers: What we know and what we don't Wendy Cozen, USC Norris Comprehensive Cancer Center, Los Angeles, CA
mutations and poorer survival in patients with diffuse large B-cell lymphoma Christopher R. Flowers	3:50–4:05 p.m.	survivors of lymphoma
4:20–4:40 p.m. Discussion / Q&A	4:05–4:20 p.m.	mutations and poorer survival in patients with diffuse large B-cell lymphoma
	4:20–4:40 p.m.	Discussion / Q&A

Session 9: Hitting Back at MYC/BCL2 Lymphomas

Moderator: David M. Weinstock, Dana-Farber Cancer Institute, Boston, MA 3:30–4:40 p.m.

Channel 2

3:30–3:35 p.m. Moderator Introduction
 3:35–3:50 p.m. Targeting MYC deregulation in cancer
 Dinis Calado, The Francis Crick Institute, London, United Kingdom

3:50–4:05 p.m.	Modeling and targeting double-hit lymphoma Bruno Amati, European Institute of Oncology, Milan, Italy
4:05–4:20 p.m.	Targeting BCL-2 with venetoclax Jennifer R. Brown, Dana-Farber Cancer Institute, Boston, MA
4:20–4:40 p.m.	Discussion / Q&A

Wednesday, August 19

Session 10: Dissecting Lymphoma Cell by Cell: A Dynamic Tale

Moderator: Kojo Elenitoba-Johnson, University of Pennsylvania School of Medicine, Philadelphia, PA 10-11:35 a.m. Channel 1

10–10:05 a.m.	Moderator Introduction
10:05–10:20 a.m.	Single-cell delineation of clonal variation and cellular tumor microenvironmental interactions in follicular lymphoma Hanlee P. Ji, Stanford University, Stanford, CA
10:20–10:35 a.m.	Pathology from the atomic scale on up Garry P. Nolan, Stanford Hospital, Stanford, CA
10:35–10:50 a.m.	Follicular lymphoma dynamics through single-cell analysis Bertrand Nadel, Centre d'Immunologie de Marseille-Luminy, Marseille, France
10:50–11:05 a.m.	Characterizing the tumor ecosystem in follicular lymphoma by CyTOF Andrew P. Weng, BC Cancer, Vancouver, BC, Canada
11:05–11:35 a.m.	Discussion / Q&A

Break

11:35-11:50 a.m.

Session 11: Can We Find and Destroy Lymphoma Precursor Cells?

Moderator: Michael R. Green, The University of Texas MD Anderson Cancer Center, Houston, TX 11:50 a.m.-1:25 p.m.

Channel 1

11:50-11:55 a.m. **Moderator Introduction**

11:55 a.m.-12:10 p.m. Tracking precursor stages of follicular lymphoma development

	Sandrine Roulland, Centre d'Immunologie de Marseille-Luminy, Marseille, France
12:10–12:25 p.m.	Defining lymphoma reservoirs: Clues from the genomics Jessica Okosun, Barts Cancer Institute, Queen Mary University of London, London, United Kingdom
12:25–12:40 p.m.	Investigating malignant transformation in Waldenstrom's Macroglobulinemia Zachary R. Hunter, Dana-Farber Cancer Institute, Boston, MA
12:40–12:55 p.m.	Defining precursors in low-grade lymphomas that seed second lymphomas David M. Weinstock, Dana-Farber Cancer Institute, Boston, MA
12:55–1:25 p.m.	Discussion / Q&A

Break

1:25-1:40 p.m.

Session 12: Using Liquid Biopsies for Real-Time Monitoring of Lymphoma Biology and Clinical Decision-Making

Moderator: David W. Scott, BC Cancer, Vancouver, BC, Canada 1:40–3:15 p.m.
Channel 1

1:40–1:45 p.m.	Moderator Introduction
1:45–2:00 p.m.	Approaches for personalized medicine in lymphoma through liquid biopsies David M. Kurtz, Stanford University, Stanford, CA
2:00–2:15 p.m.	Detecting and quantifying mutations associated with treatment resistance in aggressive lymphomas using ctDNA Ryan D. Morin, Simon Fraser University, Burnaby, BC, Canada
2:15–2:30 p.m.	Qualification and quantification of lymphoma under therapy by ctDNA analysis Davide Rossi, Institute of Oncology Research, Bellinzona, Switzerland
2:30–2:45 p.m.	ctDNA in indolent lymphoid cancers treated with targeted therapies Constantine S. Tam, St. Vincent's Hospital Sydney, East Melbourne, Australia
2:45–3:15 p.m.	Discussion / Q&A

Break

3:15-3:30 p.m.

Session 13: Preclinical Models: How to Best Predict Therapeutic Response

Moderator: Teresa Palomero, Columbia University, New York, NY

3:30–4:55 p.m. Channel 1

3:30–3:35 p.m.	Moderator Introduction
3:35–3:50 p.m.	Developing preclinical models for peripheral T-cell lymphomas Teresa Palomero
3:50–4:05 p.m.	Molecular and genetic profiling in canine lymphoma unravels targets for the human counterpart Luca Aresu, University of Turin, Turin, Italy
4:05–4:20 p.m.	Patient-derived tumor xenograft of human lymphomas Giorgio Inghirami, Weill Cornell Medicine, New York, NY
4:20–4:35 p.m.	Lymphoma organoids to study cooperative signaling and therapeutic resistance in human DLBCLs Ankur Singh, Georgia Institute of Technology, Atlanta, GA
4:35–4:55 p.m.	Discussion / Q&A

Session 14: Understanding and Targeting Aberrant Receptor Signaling in Lymphoma *Moderator: Louis M. Staudt, National Cancer Institute, Bethesda, MD*

Moderator: Louis M. Staudt, National Cancer Institute, Bethesda, MD 3:30–4:40 p.m.

Channel 2

3:30–3:35 p.m.	Moderator Introduction
3:35–3:50 p.m.	MALT1 targeting for B-cell lymphomas Lorena Fontan, Weill Cornell Medical College, New York, NY
3:50–4:05 p.m.	Tracking MYC-lymphoma evolution in vivo: Role of the B-cell receptor Stefano Casola, The FIRC Institute of Molecular Oncology (IFOM), Milan, Italy
4:05–4:20 p.m.	B-cell receptor– and co-receptor–driven metabolism reprogramming Benjamin Gewurz, Brigham & Women's Hospital, Boston, MA
4:20-4:40 p.m.	Discussion / Q&A

Break

4:55-5:10 p.m.

Keynote Address and Closing Remarks

5:10–6 p.m. Channel 1 5:10 p.m. Keynote Introduction
Ari M. Melnick, Weill Cornell Medical College, New York, NY

5:10–5:35 p.m. Cellular and molecular control of germinal center B-cell selection
Gabriel Victora, The Rockefeller University, New York, NY

5:35–5:50 p.m. Discussion / Q&A

5:50–6:00 p.m. Closing Remarks