

Sunday, December 3, 2017

5:00 p.m.–6:00 p.m.

Welcome and Opening Keynote

Grand Ballroom ABC

Fusion oncoproteins in pediatric cancer: Mechanisms and therapeutic opportunities

Scott A. Armstrong, Dana-Farber Cancer Institute, Boston, MA

6:00 p.m.–8:00 p.m.

Welcome Reception

Grand Ballroom E

Monday, December 4, 2017

7:00 a.m.–8:00 a.m.

Breakfast

Grand Ballroom D

8:00 a.m.–10:00 a.m.

Plenary Session 1: Fusion Oncoproteins as Drivers of Childhood Cancer

Grand Ballroom ABC

Session Chair: Olivier Delattre, Institut Curie, Paris, France

Identity shifts in pediatric cancers

Olivier Delattre

Subgroup-specific enhancer hijacking in medulloblastoma

Paul A. Northcott, St. Jude Children's Research Hospital, Memphis, TN

Identification of new vulnerabilities in fusion-driven pediatric cancers with functional and chemical genomic approaches

Kimberly Stegmaier, Dana-Farber Cancer Institute, Boston, MA

PAX3-FOXO1 transgenic zebrafish models identify HES3 as a mediator of tumorigenesis*

Genevieve Kendall, University of Texas Southwestern Medical Center, Dallas, TX

Undifferentiated sarcomas harbor frequent oncogenic fusions and prognostic copy number alterations: A report from the Children's Oncology Group (COG) Soft Tissue Sarcoma Committee*

Theodore Laetsch, University of Texas Southwestern Medical Center, Dallas, TX

**Short talks from proffered abstracts*

CONFERENCE PROGRAM

10:00 a.m.–10:30 a.m.

Break

Pre-Function Lobby

10:30 a.m.–12:30 p.m.

Plenary Session 2: Immunotherapy 1

Grand Ballroom ABC

Session Chair: Michael C. Jensen, Seattle Children's Research Institute, Seattle, WA

CAR T cells in the pediatric cancer clinic: Innovation and opportunities

Michael C. Jensen

The role of hypermutation and replication repair deficiency in response of childhood cancers to immune checkpoint inhibitors

Uri Y. Tabori, The Hospital for Sick Children, Toronto, ON, Canada

Engineering T cells for the immunotherapy for pediatric solid tumors

Stephen Gottschalk, St. Jude Children's Research Hospital, Memphis, TN

Analysis of immune infiltrate identifies checkpoint blockade and TLR3 activation as efficient synergistic combination of immunotherapy in rhabdoid tumors*

Amaury Leruste, Paris-Sciences-Lettres, Institut Curie Research Center, Paris, France

GD2-directed chimeric antigen receptor T cells mediate potent antitumor effect and cure in xenograft models of diffuse intrinsic pontine glioma*

Robbie G. Majzner, Stanford University School of Medicine, Palo Alto, CA

12:30 p.m.–3:00 p.m.

Lunch and Poster Session A

Grand Ballroom D&E

3:00 p.m.–5:00 p.m.

Plenary Session 3: Resistance and Refractory/Relapse

Grand Ballroom ABC

Session Chair: Yael P. Mossé, Children's Hospital of Philadelphia, Philadelphia, PA

Epigenetic mechanisms of resistance to targeted therapy

Rani E. George, Dana-Farber Cancer Institute, Boston, MA

Title to be announced

Yael P. Mossé

Neuroblastoma is a biphasic tumor

Rogier Versteeg, Academic Medical Center-University of Amsterdam, Amsterdam, The Netherlands

Genomic analysis of osteosarcoma reveals opportunities for targeted therapy*

Alejandro Sweet-Cordero, University of California San Francisco, San Francisco, CA

High-throughput drug screening identifies pazopanib and clofilium tosylate as promising treatments for malignant rhabdoid tumors*

Céline Chauvin, Paris-Sciences-Lettres, Institut Curie Research Center, Paris, France

**Short talks from proffered abstracts*

5:00 p.m.–5:15 p.m.

Break
Pre-Function Lobby

5:15 p.m.–6:30 p.m.

Policy Session: Pediatric Cancer Drug Development: Challenges and Opportunities for Success

Grand Ballroom ABC

Session Chair: Michael P. Link, Stanford University, Palo Alto, CA
(Not CME accredited)

Panelists

Peter C. Adamson, Children’s Hospital of Philadelphia, Philadelphia, PA

Hubert N. Caron, Roche Applied Science, Basel, Switzerland

Nancy F. Goodman, Kids v. Cancer, Washington, DC

Danielle Leach, St. Baldrick’s Foundation, Monrovia, CA

Michael P. Link, Stanford University, Palo Alto, CA

John M. Maris, Children’s Hospital of Philadelphia, Philadelphia, PA

Gregory H. Reaman, Food and Drug Administration, Silver Spring, MD

Tuesday, December 5, 2017

7:00 a.m.–8:00 a.m.

Breakfast
Grand Ballroom D

8:00 a.m.–10:00 a.m.

Plenary Session 4: Genomics and Clinical Trials

Grand Ballroom ABC

Session Chair: Charles G. Mullighan, St. Jude Children’s Research Hospital, Memphis, TN

Pediatric precision oncology programs in Germany and Europe

Stefan M. Pfister, German Cancer Research Center, Heidelberg, Germany

Modeling and targeting CREBBP mutations in relapsed acute lymphoblastic leukemia

Charles G. Mullighan

Implementing and evaluating precision oncology approaches for childhood cancer patients

D. William Parsons, Baylor College of Medicine, Dan L. Duncan Cancer Center, Houston, TX

A pediatric phase 1 study of larotrectinib, a highly selective inhibitor of the tropomyosin receptor kinase (TRK) family: An updated analysis*

Brian Turpin, Cincinnati Children’s Hospital Medical Center, Cincinnati, OH

Overexpression and mutations of CXorf67 in “infant-type” posterior fossa type-A ependymomas*

David Ellison, St. Jude Children’s Research Hospital, Memphis, TN

*Short talks from proffered abstracts

CONFERENCE PROGRAM

10:00 a.m.–10:30 a.m.

Break

Pre-Function Lobby

10:30 a.m.–12:30 p.m.

Plenary Session 5: Epigenetics

Grand Ballroom ABC

Session Chair: Nada Jabado, McGill University, Montreal, QC, Canada

Look at the stars and make a wish

Nada Jabado

SWI/SNF (BAF) chromatin remodeling complexes are frequently mutated in cancer: Mechanisms and vulnerabilities

Charles W. M. Roberts, St. Jude Children's Research Hospital, Memphis, TN

Mammalian SWI/SNF (BAF) complexes in pediatric cancers: New mechanisms and therapeutic opportunities

Cigall Kadoch, Dana-Farber Cancer Institute, Boston, MA

Cancer-specific retargeting of BAF complexes by a prion-like domain*

Gaylor Boulay, Massachusetts General Hospital Molecular Pathology Unit and Cancer Center, Harvard Medical School, Boston, MA

Selective gene dependencies in MYCN-amplified neuroblastoma include the core transcriptional regulatory circuitry*

Adam D. Durbin, Dana-Farber Cancer Institute, Boston, MA

12:30 p.m.–2:30 p.m.

Lunch on Own

2:30 p.m.–4:30 p.m.

Plenary Session 6: Cancer Modeling

Grand Ballroom ABC

Session Chair: Michael A. Dyer, St. Jude Children's Research Hospital, Memphis, TN

Elucidating oncohistone function using in vivo models of diffuse intrinsic pontine glioma

Suzanne J. Baker, St. Jude Children's Research Hospital, Memphis, TN

MYC activation through enhancer hijacking or focal enhancer amplification drives a subset of high-risk pediatric neuroblastoma

A. Thomas Look, Dana-Farber Cancer Institute, Boston, MA

MYCN amplification and ATRX mutations are incompatible in neuroblastoma

Michael A. Dyer

Synthetic lethality between ATRX mutations and MYCN amplification in neuroblastoma*

Maged Zeineldin, St. Jude Children's Research Hospital, Memphis, TN

**Short talks from proffered abstracts*

Hedgehog pathway drives fusion-negative rhabdomyosarcoma initiated from nonmyogenic endothelial progenitors*

Mark E. Hatley, St. Jude Children's Research Hospital, Memphis, TN

4:30 p.m.–7:00 p.m.

Reception and Poster Session B

Grand Ballroom D&E

Wednesday, December 6, 2017

7:00 a.m.–8:00 a.m.

Breakfast

Grand Ballroom D

8:00 a.m.–10:00 a.m.

Plenary Session 7: Immunotherapy 2

Grand Ballroom ABC

Session Chair: Crystal L. Mackall, Stanford University School of Medicine, Stanford, CA

Engineered T cells for pediatric cancers

Crystal L. Mackall

DNA-demethylating agents enhance cytolytic activity of CD8+ T cells and antitumor immunity

Daniel Diniz De Carvalho, Princess Margaret Cancer Centre, Toronto, ON, Canada

Strategies for harnessing both innate and adoptive immunity in hematopoietic stem cell transplantation for children with hematologic malignancies

Franco Locatelli, University of Pavia, Rome, Italy

Targeting p-TEFb as well as the EWS-ETS-specific transcriptional program in Ewing sarcoma synergistically blocks tumor growth*

Günther Richter, Technische Universität München, Children's Cancer Research Center, Munich, Germany

Mutations in microRNA processing genes dysregulate a miR-16/34-PLAG1-IGF2 axis in Wilms tumors*

Kenneth Chen, University of Texas Southwestern Medical Center, Dallas, TX

10:00–10:30 a.m.

Break

Pre-Function Lobby

**Short talks from proffered abstracts*

CONFERENCE PROGRAM

10:30 a.m. -12:30 p.m.

Plenary Session 8: New Technologies for Understanding Cancer

Grand Ballroom ABC

Session Chair: Annie Huang, The Hospital for Sick Children, Toronto, ON, Canada

Metabolic complexity in cancer cells and tumors

Ralph J. DeBerardinis, University of Texas Southwestern Medical Center, Dallas, TX

Deep epigenomic interrogation of rare pediatric brain cancers

Annie Huang

Functional proteomics and site-specific genomic rearrangements of pediatric cancers

Alex Kentsis, Memorial Sloan Kettering Cancer Center, New York, NY

Comprehensive analysis of hypermutation in human cancer reveals drivers that can be traced to the germline*

Adam Shlien, The Hospital for Sick Children, Toronto, ON, Canada

Whole-genome analysis of the epigenetic mark 5-hydroxymethylcytosine reveals differential profiles in low-, intermediate-, and high-risk neuroblastomas*

Mark Applebaum, University of Chicago, Chicago, IL

12:30 p.m.

Closing Remarks

**Short talks from proffered abstracts*