

# Conference Program

## Thursday, September 24

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**7:00 p.m.-8:00 p.m.      Opening Keynote Session**  
Ellington Ballroom  
*Session Chairperson: Charles W.M. Roberts, St. Jude Children's Research Hospital, Memphis, TN*

**Welcome remarks**  
Charles W.M. Roberts

*Keynote Address*

**Epigenetic mechanisms of tumor initiation and evolution**  
Bradley E. Bernstein, Massachusetts General Hospital, Harvard Medical School, Boston, MA

**8:00 p.m.-9:30 p.m.      Welcome Reception**  
Overlook

## Friday, September 25

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**7:00 a.m.-8:00 a.m.      Continental Breakfast**  
Terrace and Ellington Prefunction

**8:00 a.m.-10:00 a.m.      Plenary Session 1: Epigenetic Regulation of Transcription Initiation**  
Ellington Ballroom  
*Session Chairperson: Peter A. Jones, Van Andel Research Institute, Grand Rapids, MI*

**8:00 a.m.                  Genomic organization of yeast chromatin and gene regulatory complexes**  
B. Franklin Pugh, Penn State University, University Park, PA

**8:30 a.m.                  Elucidating and targeting dichotomous roles of the PRC2 complex in cancer**  
Karen M. Cichowski, Brigham and Women's Hospital, Boston, MA

**9:00 a.m.                  How DNA methylation organizes the cancer epigenome**  
Peter A. Jones

**9:30 a.m.                  Structural mechanism of sequence-specific 5-methylcytosine (5mC) recognition by AP-1 transcription factors\***  
Samuel Hong, Emory University, Atlanta, GA

**9:45 a.m.                  Transcriptional regulation mediated by biochemically distinct forms of SWI/SNF\***  
Jesse R. Raab, University of North Carolina at Chapel Hill, Chapel Hill, NC

\*Short talks from proffered papers

- 10:00 a.m.-10:30 a.m. Break**  
Ellington Prefunction
- 10:30 a.m.-12:30 p.m. Plenary Session 2: Enhancers, Epigenetics, and Cancer**  
Ellington Ballroom  
*Session Chairperson: Charles W.M. Roberts, St. Jude Children's Research Hospital, Memphis, TN*
- 10:30 a.m. Genome-wide views into the mechanisms of transcription regulation**  
John T. Lis, Cornell University, Ithaca, NY
- 11:00 a.m. SWI/SNF chromatin remodeling complex mutations in cancer: Mechanisms and potential therapeutic insights**  
Charles W.M. Roberts
- 11:30 a.m. Decoding the cancer regulome**  
John Stamatoyannopoulos, University of Washington, Seattle, WA
- 12:00 p.m. Exploring the link between Kras and histone acetylation\***  
Alessandro Carrer, Abramson Family Cancer Research Institute, Philadelphia, PA
- 12:15 p.m. Targeting super-enhancer driven oncogene transcription through cyclin-dependent kinase inhibitors\***  
Rani E. George, Dana-Farber Cancer Institute, Boston, MA
- 12:30 p.m.-2:30 p.m. Poster Session A with Lunch**  
Overlook
- 2:30 p.m.-4:30 p.m. Plenary Session 3: Control of Transcription and Elongation**  
Ellington Ballroom  
*Session Chairperson: Karen Adelman, Laboratory of Epigenetics and Stem Cell Biology, National Institute of Environmental Health Sciences, Research Triangle Park, NC*
- 2:30 p.m. An epigenome perspective of human tumor evolution**  
Joseph F. Costello, UCSF Helen Diller Family Comprehensive Cancer Center, San Francisco, CA
- 3:00 p.m. Nucleosome barriers to transcription**  
Steven Henikoff, Fred Hutchinson Cancer Research Center, Seattle, WA
- 3:30 p.m. Regulating transcription elongation at stimulus responsive genes**  
Karen Adelman
- 4:00 p.m. Using epigenetic profiling and CRISPR screens to understand cancer progression**  
X. Shirley Liu, Dana-Farber Cancer Institute, Boston, MA
- 4:30 p.m.- Evening on Own**

\*Short talks from proffered papers

## Saturday, September 26

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- 7:00 a.m.-8:00 a.m. Continental Breakfast**  
Terrace and Ellington Prefunction
- 8:00 a.m.-10:00 a.m. Plenary Session 4: Chromatin Organization**  
Ellington Ballroom  
*Session Chairperson: Sharon Y.R. Dent, The University of Texas MD Anderson Cancer Center, Smithville, TX*
- 8:00 a.m. Long-range gene regulation in the context of chromatin domains**  
Job Dekker, University of Massachusetts Medical School, Worcester, MA
- 8:30 a.m. Structure and function of BAF complexes in human cancer**  
Cigall Kadoch, Dana-Farber Cancer Institute, Boston, MA
- 9:00 a.m. A SAGA of GCN5 and USP22 in development and disease**  
Sharon Y.R. Dent
- 9:30 a.m. Mechanisms of ATP-dependent chromatin remodeling**  
Geeta Narlikar, University of California at San Francisco, San Francisco, CA
- 10:00 a.m.-10:30 a.m. Break**  
Ellington Prefunction
- 10:30 a.m.-12:30 p.m. Plenary Session 5: ncRNAs and Cancer**  
Ellington Ballroom  
*Session Chairperson: Ramin Shiekhattar, University of Miami Miller School of Medicine, Miami, FL*
- 10:30 a.m. The impact of cohesin mutations in tumor initiation: One ring, many functions**  
Iannis Aifantis, HHMI/NYU School of Medicine, New York, NY
- 11:00 a.m. Biogenesis and mechanism of action of enhancer RNAs**  
Ramin Shiekhattar
- 11:30 a.m. Functional characterization of the tumor suppressor lysine-specific methyltransferase KMT2D in lymphoma**  
Hans-Guido Wendel, Memorial Sloan Kettering Cancer Center, New York, NY
- 12:00 p.m. “Viral mimicry” as a mechanism of action for DNA-demethylating agents\***  
Daniel De Carvalho, Princess Margaret Cancer Centre, Toronto, ON, Canada
- 12:15 p.m. Analysis of enhancer transcription reveals novel gene regulatory networks in breast cancer\***  
Hector L. Franco, The Cecil H. and Ida Green Center for Reproductive Biology Sciences, UT Southwestern Medical Center, Dallas, TX
- 12:30 p.m.-2:30 p.m. Lunch on own/Free Time**

\*Short talks from proffered papers

- 2:30 p.m.-4:30 p.m. Plenary Session 6: Cancer Genomics and Epigenomics**  
Ellington Ballroom  
*Session Chairperson: Suzanne J. Baker, St. Jude Children's Research Hospital, Memphis, TN*
- 2:30 p.m. Integrative epigenomic analysis across cancer types**  
Peter W. Laird, Van Andel Research Institute, Grand Rapids, MI
- 3:00 p.m. The oncogenic role of histone H3 mutations in pediatric high-grade gliomas**  
Suzanne J. Baker
- 3:30 p.m. The cell-type specific effect of epigenomic features on cancer mutation**  
Shamil Sunyaev, Brigham and Women's Hospital, Harvard Medical School, Boston, MA
- 4:00 p.m. The oncogenic BRD4-NUT chromatin regulator drives aberrant transcription within large topological domains\***  
Erica M. Walsh, Brigham and Women's Hospital, Harvard Medical School, Boston, MA
- 4:15 p.m. GC skew defines distinct RNA polymerase pause sites in CpG island promoters\***  
Joshua S.K. Bell, Emory University, Atlanta, GA
- 4:30 p.m.-6:30 p.m. Poster Session B and Reception**  
Overlook
- 6:30 p.m.- Evening on Own**

\*Short talks from proffered papers

## Sunday, September 27

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- 7:00 a.m.-8:00 a.m. Continental Breakfast**  
Terrace and Ellington Prefunction
- 8:00 a.m.-10:00 a.m. Plenary Session 7: Epigenetic Cancer Therapies 1**  
Ellington Ballroom  
*Session Chairperson: Stephen B. Baylin, Johns Hopkins University School of Medicine, Baltimore, MD*
- 8:00 a.m. Targeting DNA methylation abnormalities as a cancer therapy strategy**  
Stephen B. Baylin
- 8:30 a.m. Multiple targetable pathways for epigenetic therapy**  
Jean-Pierre Issa, Fels Institute for Cancer Research and Molecular Biology, Temple University, Philadelphia, PA
- 9:00 a.m. Discovery of novel epigenetic targets**  
Frank Stegmeier, Novartis, Cambridge, MA
- 9:30 a.m. EZH2 inhibitors reveal broad EZH2 dependencies in multiple myeloma\***  
Shilipi Arora, Constellation Pharmaceuticals, Cambridge, MA  
*This talk is not not accredited for CME credit to permit the free flow of information of the commercial interest employees participating.*
- 9:45 a.m. A DNA hypomethylation signature predicts novel anti-tumor activity of LSD1 inhibition in SCLC\***  
Helai Mohammad, GlaxoSmithKline, Collegeville, PA  
*This talk is not not accredited for CME credit to permit the free flow of information of the commercial interest employees participating.*
- 10:00 a.m.-10:15 a.m. Break**  
Ellington Prefunction

\*Short talks from proffered papers

**10:15 a.m.-12:15 p.m. Plenary Session 8: Epigenetic Cancer Therapies 2**  
Ellington Ballroom  
*Session Chairperson: Ari M. Melnick, Weill Cornell Medical College of Cornell University, New York, NY*

**10:15 a.m. Epigenetics switches enabling transformation of B cells**  
Ari M. Melnick

**10:45 a.m. Next-generation bromodomain inhibitors**  
James E. Bradner, Dana-Farber Cancer Institute, Boston, MA

**11:15 a.m. Tazemetostat, an EZH2 Inhibitor and potential therapeutic for non-Hodgkin lymphoma**  
Jesse J. Smith, Epizyme, Inc., Waltham, MA

**11:45 a.m. A bromodomain cassette exchange strategy establishes that on-target chemical inhibition of BRD9 limits leukemia cell proliferation\***  
Anja F. Hohmann, Cold Spring Harbor Laboratory, Cold Spring Harbor, NY

**12:00 p.m. MED12 methylation by CARM1 sensitizes human breast cancer cells to chemotherapy drugs\***  
Wei Xu, University of Wisconsin, Madison, WI

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