

## 2026 A HAT Vision: 30 years of Epigenetics from Discovery to Therapy

### Schedule

March 21-23, 2026

Tsinghua University · Beijing

<b>20<sup>th</sup> March, Friday</b>	
02:00 p.m. – 04:00 p.m.	Registration and Poster set-up
<b>21<sup>st</sup> March, Saturday</b>	
08:30 a.m. – 09:00 a.m.	Opening remarks: Hongwei Wang, Yang Shi, Judith Nicholson Host: Haitao Li
<b>Session 1, Histone Modification Biology – Molecular Players and Their Functions</b> Host: Xudong Wu, Qing Li	
09:00 a.m. – 09:30 a.m.	<b>Kristian HELIN</b> , University of Copenhagen Histone Methylation and the Histone Code
09:30 a.m. – 10:00 a.m.	<b>Bing ZHU</b> , Institute of Biophysics, CAS Regulation of heterochromatin
10:00 a.m. – 10:30 a.m.	<b>Jiemin WONG</b> , East China Normal University A conserved H3K14ub-driven H3K9 methylation pathway for heterochromatin formation and inheritance
10:30 a.m. – 11:00 a.m.	<i>Group photo &amp; Break</i>
11:00 a.m. – 11:30 a.m.	<b>Yali DOU</b> , University of Southern California KMT2 Family Enzyme in Cancer
11:30 a.m. – 12:00 p.m.	<b>Yujiang Geno SHI</b> , Fudan University Taking LSD1 to a new high: LSD2 in Transcriptional Regulation and Cancer Immunotherapy
12:00 p.m. – 02:00 p.m.	<i>Lunch &amp; Break</i>
<b>Session 2, Chromatin Architecture and Structural Epigenetics</b> Host: Guohong Li, Zhucheng Chen	

02:00 p.m. – 02:30 p.m.	<b>Ming-Ming ZHOU</b> , Icahn School of Medicine at Mount Sinai Epigenetics Through a Bromodomain-Biased Lens: Three Decades from Discovery to Therapy
02:30 p.m. – 03:00 p.m.	<b>Ruiming XU</b> , Institute of Biophysics, CAS Regulation of histone acetylation: from mechanisms to applications
03:00 p.m. – 03:30 p.m.	<b>Alexander RUTHENBURG</b> , University of Chicago Promiscuous RNA binding by WDR5 remodels the KMT2A (MLL1) histone methyltransferase complex to an inactive state
03:30 p.m. – 04:00 p.m.	<i>Poster session &amp; Break</i>
04:00 p.m. – 04:30 p.m.	<b>Yael DAVID</b> , Memorial Sloan Kettering Cancer Center From mechanism to medicine: Decoding Chromatin and Cell Fate with Chemical Precision
04:30 p.m. – 05:00 p.m.	<b>Matthew SIMON</b> , Yale University A chemists focus on histone tails and the discovery of acetyl-methyllysine
05:00 p.m. – 05:30 p.m.	<b>Jaehoon KIM</b> , KAIST Transcriptional Regulation by KMT2 Family Proteins
<b>22<sup>nd</sup> March, Sunday</b>	
<b>Session 3, The Allis Vision – Shaping the Epigenetic Landscape from GCN5 to Today</b> Host: Andrew Xiao, Qiaoran Xi	
08:30 a.m. – 08:45 a.m.	<b>In Honor of Allis: Celebrating a Legacy of Epigenetic Milestones</b> (Memorial Video Screening & Group Photo)
08:45 a.m. – 09:15 a.m.	<b>Brian D. STRAHL</b> , University of North Carolina at Chapel Hill Every Amino Acid Matters: Allis’s GCN5 Legacy and the Evolving Language of Histone Modifications
09:15 a.m. – 09:30 a.m.	<b>Yifan LIU</b> , University of Southern California The Allis legacy in ciliate epigenetics
09:30 a.m. – 09:45 a.m.	<b>Jerry WORKMAN</b> , Stowers Institute for Medical Research Allis’s Footprints in Chromatin and Transcription Regulation

09:45 a.m. – 10:00 a.m.	<b>Dinshaw J. PATEL</b> , Memorial Sloan Kettering Cancer Center Structural Epigenetics Unlocked—Allis’s Insights and Collaborative Triumphs
10:00 a.m. – 10:15 a.m.	<b>Xuetong SHEN</b> , Shenzhen Bay Laboratory From Allis’s Histone Code to Actin Code
10:15 a.m. – 10:30 a.m.	<b>Haitao LI</b> , Tsinghua University The Allis Code: Shaping the Epigenetic Landscape
10:30 a.m. – 11:00 a.m.	<i>Poster session &amp; Break</i>
11:00 a.m. – 11:30 a.m.	<b>Shelley BERGER</b> , University of Pennsylvania Remembrance and Reinvention: Epigenetic and neuropeptide (re)programming of ant behavior
11:30 a.m. – 12:00 p.m.	<b>Nada JABADO</b> , McGill University An Allis Tale: The oncohistone Chronicles – Every Amino Acid Matters
12:00 p.m. – 02:00 p.m.	<i>Lunch &amp; Break</i>
<b>Session 4, Epigenetic Regulation in Development and Disease</b>	
Host: Bing Li, Mo Chen	
02:00 p.m. – 02:30 p.m.	<b>Andrew XIAO</b> , Yale University Histone variants in early development, cancer and evolution
02:30 p.m. – 03:00 p.m.	<b>Xiaohua SHEN</b> , Tsinghua University Pol II CTD canalizes genome transcription to constrain developmental plasticity
03:00 p.m. – 03:30 p.m.	<b>Wei XIE</b> , Tsinghua University Establishing the epigenome when life begins
03:30 p.m. – 04:00 p.m.	<i>Poster Session &amp; Break</i>
04:00 p.m. – 04:30 p.m.	<b>Xiaofeng CAO</b> , Institute of Genetics and Developmental Biology, CAS Epigenetic Adaptation in Plants: From Acquired Inheritance to Food Security
04:30 p.m. – 05:00 p.m.	<b>Greg WANG</b> , Duke University Chromatin dysregulation, a cancer driver

05:00 p.m. – 05:30 p.m.	<b>Petra HAJKOVA</b> , MRC Laboratory of Medical Sciences (LMS) TBD
<b>23<sup>rd</sup> March, Monday</b>	
<b>Session 5, Translational Epigenetics and Therapeutic Frontiers</b> Host: Fei Lan, Yanming Wang	
09:00 a.m. – 09:30 a.m.	<b>Or GOZANI</b> , Stanford University H3K36 Methylation in Chromatin and Cancer Biology
09:30 a.m. – 10:00 a.m.	<b>Thomas MILNE</b> , University of Oxford From Enhancer and Chromatin Biology to Precision Targeting in Leukaemia
10:00 a.m. – 10:30 a.m.	<b>François FUKS</b> , Université Libre de Bruxelles RNA Modifications in Health and Disease
10:30 a.m. – 11:00 a.m.	<i>Poster session &amp; Break</i>
11:00 a.m. – 11:30 a.m.	<b>Chao LU</b> , Columbia University Chromatin crosstalk: mechanistic insights and therapeutic opportunities
11:30 a.m. – 12:00 p.m.	<b>Yang SHI</b> , University of Oxford Chromatin regulation and cancer
12:00 p.m. – 12:10 p.m.	Closing remarks: Dinshaw J. Patel, Wei Wu