

**Sunday, March 9<sup>th</sup>**

- 3:00 pm      Check-in
- 6:00 pm      Reception (*Lobby*)
- 7:00 pm      Dinner
- 8:00 pm      Refreshments available at Bob's Pub

**NOTE:**

Meals are in the **Dining Room**

Talks are in the **Seminar Room**

Posters are in the **Lobby**

**Monday, March 10<sup>th</sup>**

All talks are 20 minutes  
plus 10 min for Q&A

7:30 am Breakfast (*service ends at 8:45am*)

**9:00 am Session 1**  
**Chair: Timothee Lionnet**

9:00 am **Wouter de Laat**, Hubrecht Institute  
*Systematic mapping reveals large-scale and dynamic association of genomic regions with pericentromeric heterochromatin*

9:30 am **Job Dekker**, University of Massachusetts Medical School  
*Chromosome folding and gene regulation*

10:00 am **Denis Duboule**, University of Geneva and EPFL Lausanne  
*A regulatory switch between topological domains during development*

10:30 am Break

**11:00 am Session 2**  
**Chair: Enrico Gratton**

11:00 am **Rob Phillips**, California Institute of Technology  
*The statistical mechanical Genome*

11:30 am **Leonid Mirny**, Massachusetts Institute of Technology  
*Polymer models of chromosome organizations*

12:00 pm **Mario Nicodemi**, University of Naples "Federico II"  
*A view from polymer physics of chromatin spatial organisation*

12:30 pm Lunch (*service ends at 1pm*)

**2:00 pm Session 3**  
**Chair: Wendy Bickmore**

2:00 pm **Joanna Wysocka**, Stanford University School of Medicine  
*Enhancers in regulation of developmental plasticity*

2:30 pm **Clarissa S. Scholes**, Harvard Medical School  
*Effects of gene locus organization on enhancer activity*

3:00 pm **Robert Hill**, University of Edinburgh  
*Regulatory landscape of the Shh gene*

## Long-Range Genome Organization and Transcription Dynamics

3:30 pm Break

**4:00 pm** **Session 4**  
**Chair: Doug Higgs**

4:00 pm **Thomas Gregor**, Princeton University  
*From single molecules to transcription dynamics in development*

4:30 pm **Michael Levine**, University of California, Berkeley  
*Pause control in development*

5:00 pm **David Stern**, Janelia Farm Research Campus/HHMI  
*How a complex cis-regulatory region encodes robustness and underlies morphological evolution*

5:30 pm Poster Reception

7:00 pm Dinner

8:00 pm Refreshments available at Bob's Pub

**Tuesday, March 11<sup>th</sup>**

- 7:30 am Breakfast (*service ends at 8:45am*)
- 9:00 am Session 5**  
**Chair: Thomas Gregor**
- 9:00 am **Kerstin Bystricky**, University of Toulouse  
*Visualisation of folding and dynamics of single gene loci*
- 9:30 am **Enrico Gratton**, University of California, Irvine  
*Nanoscale topological structure of chromatin in live cells revealed by gold-enhanced fluorescence*
- 10:00 am **Zhe Liu**, Janelia Farm Research Campus/HHMI  
*Imaging pluripotency factor target search and enhancer binding dynamics at the single-molecule level*
- 10:30 am Break
- 11:00 am Session 6**  
**Chair: Joanna Wysocka**
- 11:00 am **Wendy Bickmore**, University of Edinburgh  
*Looking small to understand large: Chromatin folding and gene regulation*
- 11:30 am **Musa Mhlanga**, Council for Scientific and Industrial Research (CSIR)  
*Chromosomal contact permits transcription between coregulated genes*
- 12:00 pm **Martha L. Bulyk**, Brigham & Women's Hospital and Harvard Medical School  
*The genomic landscape of NF- $\kappa$ B binding in lymphoblastoid B-Cells*
- 12:30 pm Lunch (*service ends at 1pm*)
- 1:15 pm Tour (*optional – meet at reception*)
- 2:15 pm Session 7**  
**Chair: James Kadonaga**
- 2:15 pm **Jie Xiao**, Johns Hopkins University  
*Probing transcription factor-mediated DNA looping in live cells at the single molecule level*

## Long-Range Genome Organization and Transcription Dynamics

- 2:45 pm **Johan Elf**, Uppsala University  
*Non-equilibrium contributions to transcription factor mediated gene regulation*
- 3:15 pm **Timothee Lionnet**, Janelia Farm Research Campus/HHMI  
*Probing mechanisms of transcription activation at the single molecule level*
- 3:45 pm Break
- 4:15 pm Session 8**  
**Chair: Martha Bulyk**
- 4:15 pm **Antoine Coulon**, National Institutes of Health  
*Kinetic competition during the transcription cycle results in both co- and post-transcriptional splicing at the same gene*
- 4:45 pm **Alberto R. Kornblihtt**, Universidad de Buenos Aires  
*Chromatin, Pol II elongation and alternative splicing*
- 5:15 pm **Daniel Zenklusen**, Université de Montréal  
*Dissecting eRNA mediated transcription regulation in single cells*
- 5:45 pm Poster Reception
- 7:15 pm Dinner
- 8:15 pm Refreshments available at Bob's Pub

**Wednesday, March 12<sup>th</sup>**

- 7:30 am Breakfast (*service ends at 8:45am*)
- 9:00 am Session 9**  
**Chair: Rob Phillips**
- 9:00 am **Doug R. Higgs**, Weatherall Institute of Molecular Medicine  
*Linking cis-acting elements and analysing how variants of such elements may influence the chromosomal landscape*
- 9:30 am **Gary Felsenfeld**, National Institutes of Health  
*CTCF function and long range interactions*
- 10:00 am **Gerd Blobel**, Children's Hospital of Philadelphia  
*Reactivation of developmentally silenced globin genes by forced chromatin looping*
- 10:30 am Break
- 11:00 am Session 10**  
**Chair: Carl Wu**
- 11:00 am **James T. Kadonaga**, University of California, San Diego  
*The TCT core promoter motif functions with a specialized transcription system*
- 11:30 am **James A. Goodrich**, University of Colorado at Boulder  
*Regulation of RNA polymerase II by noncoding RNAs*
- 12:00 pm Closing Discussion
- 12:30 pm Lunch & Departure
- 1:00 pm First shuttle to Dulles  
2:00 pm Second shuttle to Dulles  
3:00 pm Last shuttle to Dulles