

Multidimensional Genomics: The 3D/4D organization of chromatin November 13-15, 2017

Monday, 13 November 2017

8.30-9.00 Registration

9.00-9.15 Welcome by Modesto Orozco

Session I

Chair: Modesto Orozco

9.15-10.00 Steroid Receptor Binding Domains mediate receptor dependent

intra-TAD folding and global genome organization in breast cancer

cells

Miguel Beato, CRG (Barcelona, Spain)

10.00-10.45 Oncogenic enhancers dynamically associated with MYC in 3D

influence its post-transcriptional regulation by mediating

gene-nuclear pore interactions

Rolf Ohlsson, Karolinska Institutet (Stockholm, Sweden)

- 10.45-11.15 Coffee break and poster session
- 11.15-12.00 Role of Polycomb and MLL complexes in nuclear architecture and gene regulation

Luciano Di Croce, CRG/ ICREA (Barcelona, Spain)

12.00-12.20 SHORT TALK 1 Chromosome 3D organization is a driver of position

effects in Drosophila

Guillaume J. Filion, CRG (Barcelona, Spain)

12.20-13.05 Functional impact of altering chromatin accessibility on mutational

burden, topography, and tumor progression

Salvador Aznar-Benitah, IRB Barcelona/ICREA (Barcelona, Spain)

13.05-15.05 Lunch (FrescCo, C/ Carme, 16)

Session II

Chair: Giacomo Cavalli

15.05-15.50 Chromosome folding by loop extrusion

Leonid Mirny, Massachusetts Institute of Technology (Cambridge, MA, USA)

15.50-16.10 SHORT TALK 2 PoSTAC (Polycistronic SunTAg modified CRISPR) enables live-cell and super-resolution visualization of multiple genes

Maria Victoria Neguembor, CRG (Barcelona, Spain)

16.10-16.55 New technologies to understand the link between genome structure and activity

Jim R. Hughes, MRC Weatherall Institute of Molecular Medicine (Oxford, UK)

- 16.55-17.25 Coffee break and poster session
- 17.25-17.45 SHORT TALK 3 Network methods uncover the 3D organization of replication factories

Vera Pancaldi, Barcelona Supercomputing Centre (Barcelona, Spain)

17.45-18.30 Analysis of Chromatin Organization after Acute Loss of CTCF in Mouse Embryonic Stem Cells

Bing Ren, University of California San Diego (La Jolla, CA, USA)

18.30 End of session

Tuesday, 14 November 2017

Session III

Chair: Marc Martí-Renom

9.00-9.45 A quantitative analysis of the hierarchical folding of mammalian

chromosomes

Luca Giorgetti, Friedrich Miescher Institute for Biomedical Research

(Basel, Switzerland)

9.45-10.30 Multiscale modeling of the chromatin fiber: from nucleosomes to

chromosomes

Tamar Schlick, Courant Institute of Mathematical Sciences (New York, USA)

- 10.30-11.00 Coffee break and poster session
- 11.00-11.45 Epigenome folding: a main driver of 3D chromatin organization
 Daniel Jost, University Grenoble Alpes (Grenoble, França)

- 11.45-12.05 SHORT TALK 4 Computational modeling reveals 4D genome reorganization during stem cell differentiation

 Jonas Paulsen, University of Oslo Faculty of Medicine (Oslo, Norway)
- 12.05-12.50 Hi-C-constrained physical models of human chromosomes recover functionally-related properties of genome organization

 Cristian Micheletti, International School for Advanced Studies (Trieste, Italy)
- 12.50-15.00 Lunch (FrescCo, C/ Carme, 16)

Session IV

Chair: Giacomo Cavalli

- 15.00-15.45 Cracking the Nucleus-Visualizing the Higher Order Coding Structures of DNA
 Clodagh O'Shea, Salk Institute for Biological Studies (La Jolla, CA, USA)
- 15.45-16.05 SHORT TALK 5 Genome-wide mapping of lamin B1 reveals the existence of dynamic and functional euchromatin lamin B1 domains (eLADs)
 Sandra Peiró, Vall d'Hebron Institute of Oncology (Barcelona, Spain)
- 16.05-16.50 The nanoscale structure and looping of chromatin fibers in somatic and stem cells

 Maria Pia Cosma, CRG/ICREA (Barcelona, Spain)
- 16.50-17.20 Coffee break and poster session
- 17.20-17.40 SHORT TALK 6 Genome Architecture Mapping of Rare Cell Types in the Brain

 Warren Winick-Ng, Berlin Institute for Medical Systems Biology (Berlin, Germany)
- 17.40-18.25 Updates on imaging and homolog pairing
 Ting Wu, Harvard Medical School (Boston, MA, USA)
- 18.25 End of session
- 20:30 Speakers dinner

"Restaurant Agüelo013" (Avinyó, 37)

Wednesday, 15 November 2017

Session V

Chair: Modesto Orozco

9.00-9.45 3D Chromosome and Genome Dynamics

Peter Fraser, The Babraham Institute (Cambridge, UK)

9.45-10.30 Chromatin structure in yeast: from beads on a string to 3D folding

Oliver J. Rando, University of Massachusetts Medical School (Worcester, MA,

USA)

10.30-11.00 Coffee break and poster session

11.00-11.20 SHORT TALK 7 YY1 is a general structural component of

enhancer-promoter loops

Alicia V. Zamudio, Whitehead Institute for Biomedical Research

(Cambridge, MA, USA)

11.20-12.05 Genome architecture: mechanisms of 3D chromatin folding

Ana Pombo, MDC (Berlin, Germany)

12.05 Closing remarks