

Detailed scientific program (including keynote lectures, poster sessions & open discussions):	
DAY 1, OCTOBER 15	SUNDAY
15:00	REGISTRATION OPEN
17:45	OPENING REMARKS: Organizing Committee
18:00 – 19:00	KEYNOTE LECTURE: Tom Misteli The stochastic nature of genome organization and function
19:00 – 20:00	<i>WELCOME DRINK</i>
20:00	<i>DINNER</i>
DAY 2, OCTOBER 16	MONDAY
8:30 – 10:30	SESSION 1: CHROMATIN MODIFICATIONS Speakers: Petra Hajkova: Plasticity and dynamics of epigenetic information Simone Sidoli: Mass-spectrometric and proteomic approaches to study histone modifications Alejandra Loyola: Functions of histone modifications and histone variants Two short talks selected from the abstracts
10:30 – 11:00	<i>COFFEE BREAK</i>
11:00 – 13:00	SESSION 1 (continued): CHROMATIN MODIFICATIONS Speakers: EMBO YIP Lecture - Tuncay Baubec: High-throughput functional analysis of epigenetic gene regulatory circuits Till Bartke: Decoding chromatin states by proteomic profiling of modification-dependent nucleosome readers Andreas Ladurner: Chromatin regulation by ADP-ribosylation Two short talks selected from the abstracts
13:00 – 15:00	<i>LUNCH & FREE TIME</i>
15:00 – 17:30	SESSION 2: SPATIAL & INTERACTION PROTEOMICS Speakers: Michiel Vermeulen: Investigating the regulation of gene expression using quantitative interaction proteomics Kathryn Lilley: Spatial maps of the subcellular proteome Anne-Claude Gingras: Mapping the dynamics of cellular compartments by proximity-dependent biotinylation Two short talks selected from the abstracts Poster flash talks (for poster session 1)
17:30 – 20:00	POSTER SESSION 1
20:00	<i>DINNER</i>
DAY 3, OCTOBER 17	TUESDAY
8:30 – 10:30	SESSION 3: STRUCTURAL PROTEOMICS Speakers: Hitoshi Kurumizaka: Structural biology of chromatin Pedro Beltrao: Multi-omics data integration of post-translational modification landscapes Georg Kustatscher: Probing the dynamics of chromosome composition by computational proteomics Two short talks selected from the abstracts
10:30 – 11:00	<i>COFFEE BREAK</i>
11:00 – 12:30	SESSION 4: MULTI-OMICS DATA INTEGRATION Speakers: Alexey Nesvizhskii: FragPipe: a comprehensive computational platform for proteomics, proteogenomics, and chemoproteomics Jussi Taipale: Towards predicting gene expression from DNA sequence Two short talks selected from the abstracts
12:30 – 13:15	WOMEN IN SCIENCE LECTURE: Speaker to be confirmed Title to be confirmed
13:15 – 15:00	<i>LUNCH & FREE TIME</i>
15:00 – 17:00	PROBLEM SOLVING WORKSHOP 1:1 speaker/participant pairs

17:30	EXCURSION TO CHANIA OLD TOWN AND DINNER IN CHANIA (optional, extra cost)
20:00	<i>DINNER AT HOTEL (for those not participating in the excursion)</i>
DAY 4, OCTOBER 18	WEDNESDAY
8:30 – 10:30	SESSION 5: QUANTITATIVE APPROACHES TO TRANSCRIPTION Speakers: Tineke Lenstra: Understanding transcription in single living cells Gordon Hager: Transcription factor dynamics in live cells, and interactions with the cohesin complex Naama Barkai: How transcription factors find their binding sites in large genomes – the role of intrinsically disordered regions Two short talks selected from the abstracts
10:30 – 11:00	<i>COFFEE BREAK</i>
11:00 – 12:30	SESSION 6: HIGH-RESOLUTION IMAGING OF THE NUCLEUS Speakers: Melike Lakadamyali: Super-resolution imaging of chromatin organization and dynamics Musa Mhlanga: Visualization of chromatin dynamics by live cell imaging Two short talks selected from the abstracts
12:30 – 13:15	EMBO SCIENCE POLICY LECTURE: Speaker to be confirmed Title to be confirmed
13:15 – 15:00	<i>LUNCH & FREE TIME</i>
15:00 – 17:30	SESSION 7: CHROMATIN & METABOLISM Speakers: Kathryn Wellen: Compartmentalization of acyl-CoA metabolism and roles in chromatin regulation Axel Imhof: Cross-talk between metabolism and histone modifications Nathaniel Snyder: Compartmentalization of coenzyme metabolism within cells Two short talks selected from the abstracts Poster flash talks (for poster session 2)
17:30 – 20:00	POSTER SESSION 2
20:00	<i>DINNER</i>
DAY 5, OCTOBER 19	THURSDAY
8:30 – 10:30	SESSION 8: SINGLE CELL GENOMICS & PROTEOMICS Speakers: Jop Kind: Spatial genome organization and gene regulation in single cells Nikolai Slavov: Analyzing the nuclear proteome of single-cells Maria Colomé-Tatché – to be confirmed Two short talks selected from the abstracts
10:30 – 11:00	<i>COFFEE BREAK</i>
11:00 – 13:00	SESSION 9: DNA REPLICATION & REPAIR Speakers: Genevieve Almouzni: Maintenance of the epigenome through cell divisions Constance Alabert: Mechanisms of DNA replication in the context of chromatin Evi Soutoglou: Chromatin and nuclear dynamics in the maintenance of replication fork integrity Two short talks selected from the abstracts
13:00 – 15:00	<i>LUNCH & FREE TIME</i>
15:00 – 17:15	SESSION 10: NUCLEAR ARCHITECTURE Speakers: EMBO KEYNOTE LECTURE - Ana Pombo: Multiome-GAM: connecting cell states with 3D genome structure Nils Krietenstein: Mapping the 3D genome with nucleosome resolution Wendy Bickmore: Enhancer-promoter communication: is close enough, enough? Two short talks selected from the abstracts
17:30 – 18:15	OPEN DISCUSSION: Current and future challenges in nuclear biology
18:15 – 18:30	BEST PRESENTATION & POSTER PRIZE AWARDS

	CLOSING REMARKS: Organizing Committee
19:00	<i>DINNER</i>
20:30	<i>FAREWELL SOCIAL EVENT (Beach Party)</i>
DAY 6, OCTOBER 20	FRIDAY
	BREAKFAST AND DEPARTURE