Poste	ers A-Z
Akan, Izlem Su Molecular dissection of cofactor function in mammalian gene activation	51
Akilli, Nazli SUMOylation regulates the phase behavior and function of PRC1 in Drosophila melanogaster	52
Aktay, Serhat Mapping the functional genomic landscape de novo with divergent transcription analysis of PRO-seq data	53
AlBawardi, Waad Irregular nucleosome spacing drives chromatin into a more open and dynamic state	54
AlHaj Abed, Jumana Uncovering homologous chromosome communication in Somatic cells	55
Aloisi, Sara Unveiling non-canonical functions of CoREST complexes in neuroblastoma	56
Alshanbari, Arwa Characterisation of cyclin K-SET1A interaction to develop probes for its role in acute myeloid leukaemia	57
Ant, Luciana Chromatin landscape of endometrial tumor cells of tamoxifen users is associated with loss of progestin regulation	58
Antonovic, Milan Re-constructing the prdm14 regulatory landscape using synthetic biology	59

60

Cancelled

Arumugam, Thiruma (Garrie) Intergenerational impact of parental interest metabolic phenotypic traits	ermittent fasting on offspring	61
Baderna, Valentina A chromatin-integrated reporter assay epigenetic contribution to enhancer ac	5 5	62
Baeg, Kyungmin Tracking initiation of paraspeckle asset	mbly in real-time	63
Baehr, Stephan Basic modeling of information storage	loss in chromatin	64
Balaramane, Delphine MethyLasso: a segmentation approach patterns and identify differentially meth whole-genome datasets		65
Balwierz, Piotr ISMARA: completely automated inferer networks from high-throughput data.	Presenter: Pachkov, Mikhail nce of gene regulatory	66
Barbieri, Elisa OTX2 controls chromatin accessibility cells	in formative pluripotent stem	67
Becker, Robert Leveraging STARR-Seq to identify enhancements	ancers for AAV gene therapy	68
Bompadre, Olimpia Liebenberg syndrome severity arises for topology and ectopically transcribing of		69
Borst, Noa Transcriptional hubs as agents of robu	stness and evolvability	70

Bosnjakovic, Marta	
Analysis of transcription initiation in early embryonic development of the surface and cave forms of Astyanax mexicanus highlight regulatory rules of harsh environment adaptation	71
Boulanger, Mathias Quantifying coordinated regulation of cis-regulatory elements using long-read sequencing	72
Brennecke, Theresia Mechanism of promoter-proximal pause release: DISF and NELF phosphorylation by P-TEFb	73
Bressin, Ann Katrin Presenter: Mayer, Andreas Regulated widespread enhancer transcription as revealed by high-sensitivity nascent transcript sequencing	74
Campos Fornés, Víctor Re-wiring of KRAB zinc finger protein - transposable element regulatory networks in response to stimulation in human neurons	75
Carew, Frederic PLANT GAM discovers topologically associating domains in arabidopsis	76
Castiglione, Melissa Defining activities of KDM5 essential to development and viability	77
Chaban, Anastasiia Real-time tracking of early ribosomal RNA transcription and processing	78
Chakraborty, Chaitali Highly interconnected enhancer-promoter interactions in defining gene expression in glioblastoma	79
Chatsirisupachai, Kasit Mammalian promoters are characterised by low frequency of RNA Pol Il occupancy	80

Contreras, Aida	Presenter: Perea-Resa, Carlos	
Recycling Cohesin across Mitosis to	o re-start Transcription	81
Cooper, David Transcriptional activation domains f	Presenter: Erkine, Alex	82
hydrophobic-acidic surfactants	unction as ampinphine	02
Cordier, Julie		00
Using artificial regulatory landscape enhancer's regulatory input to two p		83
Cortez, Kaitlyn		
Understanding how GA-binding TFs gene expression	mediate 3D loop contacts and	84
Cottingham, Kendall	Presenter: Hall, Hana	
R-loop accumulation and transcripti neurodegeneration of tauopathy mo		85
Cruz-Ruiz, Samantha Functional characterization of regula	otory long non-coding DNAs in	86
response to transcriptional stress	atory long non-coding KNAS in	00
Currie, Madeline	OD annual of an in Lancau and	07
Discovery and characterization of H	2B sumoyiation in numan cells	87
Daouk, Reem		
L1 RNA re-expression through heter palmitate induced inflammatory resp		88
Dar, Mohd Saleem		
Deciphering the epigenetic roles of I HNSCC	NSD3 isoforms in HPV- negative	89
Davies, Molly		
PAX8 master transcription factor co- independently of co-regulators in ov		90
De Putter, Milan	ad BUOVO in masses and massic	0.1
Molecular mechanisms of RHOX5 ar stem cells and primordial germ cells		91

Dehingia, Bondita RNA binding proteins contribute to the consolidation CTCF-mediated chromatin architectural features upon loss of pluripotency	92
Devlin, Jennifer Control of RNA polymerase II pausing by a CDK11-CDK9 regulatory axis	93
Dittrich, Marianna SUPT5H influences cell identity in pancreatic ductal adenocarcinoma	94
Doyle, Emma Mechanistic evaluation of ASXL1 gain-of-function variants on polycomb activity in cancer and developmental syndromes	95
Dulja, Alessandro Exploring the role of transcription in topology during early Drosophila embryogenesis	96
Ehara, Haruhiko Cryo-EM analysis on the co-transcriptional nucleosome disassembly/reassembly during transcription elongation	97
Elchert, Alexandra Regulation of the histone methyltransferase Set2 by the histone chaperone Spt6	98
Espie-Caullet, Tristan Identification of novel non-canonical regulators of chromatin-mediated alternative splicing	99
Fabbro Burtschell, Leonie Function of H3K9 methylation in C. elegans developmental timing.	100
Fallon, Conrad Quantitative modulation of spatial enhancer activity as a function of transcription factor binding site affinity and orientation	101

Fasouli, Eirini Sofia Distinct role of STAT3 and STAT5 transcriptional networks in MDS to AML leukemic transformation	102
Fettweis, Gregory Transcription factors associate with genome architecture machinery to curve gene expression	103
Fischer, Martin p53 enhancer regulation reveals decoupling of transcription factor binding and transcription initiation	104
Fontalva, Sara Human prefoldin regulates gene transcription through FACT-mediated chromatin dynamics	105
Forouzanfar, Faezeh Presenter: Mendoza, Manuel Gene-specific transcript buffering revealed by perturbation of coactivator complexes	106
Gaetani, Paola Dynamics of transcription silencing during mitosis	107
Gannon, Dáire Endogenous degron tagging of p300 and CBP to investigate their roles in transcription and chromatin regulation in vivo	108
Gelder, Katie Investigating the cooperative role of IDRs on the biological phase separation of CBP	109
Geller, Merle Presenter: Liefke, Robert A paradigm for multivalent chromatin binding: protein polymerization mediated by the SAM domain and an intrinsically disordered region are both required for the full function of SAMD1 on chromatin.	1 110
Goldmann, Dora Presenter: Jafari, Shadi A high-resolution approach for detecting skewed X chromosome inactivation: implications for immunological and genetic studies in	111

Poste	rs A-Z
Goll, Sandra Molecular dissection of BAF complex assembly	112
González, Araceli Fine-tuning phosphorylation of RNA polymerase II subunit Rpb4 plays a fundamental role in gene expression	113
González-López, Silvia The dynamic establishment of enhancer function	114
Grabowska, Agnieszka The role of RNA Polymerase II condensates in the regulation of gene expression during embryonic development	115
Greulich, Franziska BRG1 and glucocorticoids – more than gene activation	116
Griffa, Alberto Single-cell multi-omic analysis reveals principles of transcription-chromatin interaction during embryonic development	117
Grimm, Christina Long-read-transcriptome-sequencing of CLL and MDS patients reveals common effects of SF3B1 mutations on RNA splicing	118
Guerrero Carreño, Xiomara Presenter: Papadopoulos, Petros Gene expression profiling of cultured proerythroblasts and mature megakaryocytes from Essential Thrombocythemia patients	119
Gupta, Himanshu Autoacetylation driven phase separation of TIP60 is crucial for its functions	120
Gutierrez Santiago, Francisco Presenter: Libri, Domenico R-loops revisited through the magnifying glass of H-CRAC: the 3'-end connection	121

Hafizi Yazdabadi, Mahtab Deciphering the molecular mechanism remodelling ability of HMGA protein us spectroscopy		122
Harvey, Zachary H2A.Z sets the transcriptional ground interaction with Spt6	state of chromatin via	123
Hauth, Antonia Escape from X inactivation is directly	Presenter: Loda, Agnese modulated by Xist RNA levels	124
Heinzl, Monika Understanding the sequence-encoding learning	g of gene regulation using deep	125
Henneman, Nathaniel Metabolic resilience in fasting depend coordinate epigenetic writers and RNA		126
Herrero Ruiz, Andrés Manuel A new promoter element for human tR	NA genes	127
Hirata, Yoshito Improving the resolution of 3D chromo on single diploid cell Hi-C data	osomal reconstructions based	128
Hisler, Vincent TBP binding and RNA polymerase II tr partially assembled TFIID complexes i		129
Hofer, Ines Haematopoietic landscape dynamics o latency.	of human cytomegalovirus	130
Hu, Yimin Bacterial histone HBb compacts DNA	by bending	131
Huynh, Duyen Effective in vivo binding energy lands transcription factor function	Presenter: Gebhardt, Christof M. cape reveals kinetic details of	132

Islam, Zubairul Enhancer mediated long range gene hubs define signaling response	133
Janowska, Aleksandra RAS-driven chromatin rewiring in slow-cycling cells	134
Javed, Aqib Epidrugs induce growth arrest and apoptosis in cancer cells via TSGs reactivation	135
Jia, Zeran ACC-seq reveals transcriptional condensates that regulate intermediate expression levels	136
Jurriens, Daphne Single molecule biology: unravelling the mechanisms underlying chromatin organization and transcription	137
Karpinska, Magdalena CTCF depletion decouples enhancer-mediated gene activation from chromatin hub formation during cellular differentiation	138
Kaur Dhall, Jasmine IP3 receptor mediated changes in transcription regulation by RNA PollI and chromatin dynamics in Drosophila brain tissue	139
Keß, Simone Dynamic regulation of developmental gene expression by enhancer RNAs during zebrafish development	140
Kirdok, Idil Novel roles for integrator complex in the context of BRCA1 deficiency	141
Klápštová, Veronika Structural characterization of the interaction between BRCA1-BARD1 and RNA polymerase II	142

Knerr, Julian Formin-mediated nuclear actin assembly at androgen receptors promotes transcriptional condensate formation for androgen signaling	143
Kolly, Dominique Presenter: Mayran, Alexandre Winner takes all or fair share? competing for shared enhancers	144
Kook, Hyun Circular RNA circSMAD4 regulates cardiac fibrosis by targeting miR-671-5p and FGFR2 in cardiac fibroblasts	145
Kopczynska, Magda Histone methyltransferase SETD2 facilitates transcription termination and pre-mRNA 3' end cleavage of fast-transcribed genes	146
Cancelled	147
Kribelbauer, Judith EXTRA-seq: a genome-integrated extended reporter assay to uncover the building principles of Enhancer-Promoter communication	148
Kuan, Chi Revealing the function of GDAC in plant immune responses under elevated temperatures	149
Kuester, Karolin Disease modelling of Weaver Syndrome in human cortical organoids	150
Kujirai, Tomoya Structural insights into transcription elongation on chromatin in cells	151
Kumar, Yatendra Unzipping the core: acetylation at the nucleosome dyad is associated with fragile nucleosomes at active promoters, enhancers, and insulators	152

Pos	ters A-Z
Lakhe, Dhanashree Attenuation of the stress response preserves genome integrity in Drosophila germ cells	153
Lee, Joo TRACK-IT: the next generation of high-speed, ultra-resolution live-microscopy of 3D genome motion across multiple genomic distances	154
Lee, Su Kyong Identifying regulators of transcriptional reinitiation with a proximity biotin labeling assay in yeast	155
Lee, Su Kyong Presenter: Rosonina, Emanuel Using biotin pulse labeling to study transcriptional dynamics	156
Leen, Eoin N-myc recruits the general transcription factor TFIIIC via the DNA binding surface of TFIIIC5	157
Lejart, Audrey Characterization of the organized structure of the chromatin in human monocytes	158
Leo, Alessandro Host deubiquitinase BAP1 supports KSHV lytic viral gene expression	159
Levesque, Mathieu Defining the cellular role of an endogenous truncated isoform of ATRX	160
Linhartová, Katerina Sequence and structural determinants of RNAPII CTD	161

Page 23

The impact of enhancer-mediated transcription on higher order

162

phase-separation and phosphorylation by CDK7

chromosome architecture: a single TAD perspective

Logie, Colin

Lone, Kaisar Unveiling the Epigenetic Influence of MLL Family Members on Ribosomal DNA Chromatin Dynamics	163
Lu, Grace Hui-Chun Intricate and highly dynamic sequence of regulatory events during the process of neural induction.	164
Luise White, Paige Presenter: Mueller, Ferenc Transcription initiation base choice defines post-transcriptional mRNA fate	165
Lundkvist, Moa Presenter: Schwartz, Yuri What are epigenetic memories good for?	166
Lux, Vanda Structural studies of JASPer protein and its complex with JIL-1 kinase to unravel preservation of active transcription	167
Magnitov, Mikhail ZNF143 is a transcriptional regulator of nuclear-encoded mitochondrial genes that acts independently of looping and CTCF	168
Majchrzycka, Blanka The promoter-proximal region mediates responsiveness to regulatory landscapes	169
Makasheva, Kristina Modular three-color live cell imaging system to study enhancer-promoter dynamics with transcriptional output	170
Maljanen, Katri Modeling effects of the epigenome on somatic structural variant formation	171
Mâmmer Bouhou, Widad Providing epigenomic single-cell sequencing technologies to the science community	172

Manjunath, Meghana Role of lysine specific demethylase-1 (KDM1A) in pancreatic ductal adenocarcinoma (PDAC)	173
Mann, Rajat RNA buffers noise in transcription factor binding and condensate formation	174
Maqsood, Khizra Presenter: Polvora Brandao, Duart Explainable artificial intelligence identifies novel mammalian enhancers and their epigenetic code	e 175
Mariani, Andrea Regulatory logic of polycomb genes is embedded in the local genomic context	176
Martianov, Igor Mouse model to study the effect of mutations in TFIID subunit TAF13 on neurodevelopment	177
Martinez Corral, Rosa Transcription factor duality in a fixed molecular context	178
Massah, Shabnam Comparing the phase separation of the androgen receptor (AR) and its splice variant (AR-V7) in prostate cancer	179
McCann, Abby Context dependent activity of p63-bound gene regulatory elements	180
Mimoso, Claudia Presenter: Vlaming, Hanneke 5' splice sites protect mRNAs from Restrictor's promiscuous activity on early elongating RNA Polymerase II	181
Mizeret, Romane Concentration-dependent regulation of genome-wide transcription factor binding	182
Montez, Miguel Cold-driven nucleosome dynamics underlying epigenetic silencing of Arabidopsis FLC	183

Monziani, Alan BAHCC1 positively regulates gene e SIN3-HDAC complex	expression by antagonizing the	184
Morao, Ana Karina DNA supercoiling modulates transc dynamics in a gene-orientation depo		185
Moreno, R. Yvette Thr4 phosphorylation primes Ser2 p regulates 3'-end processing	Presenter: Zhang, Y. Jessie phosphorylation on RNA Pol II and	186
Morillo Leonardo, Sarah Modeling single-cell chromatin acce factor activities	essibility in terms of transcription	187
Müller, Carolin Direct detection of DNA and RNA mosequencing reveals transcriptional of		188
Müller, Marisa Presenters Structural basis of histone acetylation the MSL complex	s: Doudin, Matteo; Kolesnikova, Olga on by the reader/writer module of	189
Museridze, Mariam Re-evaluating enhancer modularity approaches in Drosophila	and boundaries with quantitative	190
Nemcko, Filip Proteome-scale tagging and functio by ORFtag	onal screening in mammalian cells	191
Nenova, Aleksandra Investigating allelic imbalance in F1 strains	mice from recently diverged	192
Newland, Jake Eight principal chromatin states fun genome into developmental and ho		193

Newland, Jake Differing modes of epigenetic repression in the Drosophila brain revealed by large-scale targeted DamID profiling and chromatin state analysis	194
Nissen, Itzel Gene regulatory networks mediating neuron-to-glioma synaptic communication in glioblastoma	195
Nolan, James Presenter: Conway, Eric AEBP2-PRC2.2 subcomplex dependency in germinal centre lymphoma	196
Océane, Tourniere Dosage compensation is dispensable in most tissues but essential in the stem cells of the respiratory system for male survival	197
Olan, Ioana Redefining 3D chromatin compartments	198
Oliveros Diez, Winona Inter-individual and inter-tissue variation of DNA methylation	199
Olsen, Karl Presenter: Braun, Sigurd Shaping heterochromatin boundaries: Dynamic interplay and turnover of chromatin readers	200
Oomen, Marlies The transcriptional repertoire of transposable elements during mammalian preimplantation development	201
Oravecz, Attila Live imaging upon induced gene transcription predicts differential dynamic behavior between RNA polymerase II and initiation factors	202
Özdemir Aygenli, Bihter Functional characterization of the E3 ubiquitin ligase UHRF1	203

Paic, Antonia Developing an inducible expression system for extremophilic yeast Debaryomyces hansenii, a synthetic biology chassis compatible with hypersaline environments	204
Paldi, Flora A histone hyperacetylation pulse induces cellular memory of 3D genome folding and gene regulation	205
Palma, Flavio Oxidation of H3.1Cys96 promotes chromatin remodeling and transcriptional profile adaptation	206
Palmer, Marley Conserved short linear motif within KSHV viral protein ORF45 promotes its interaction with lytic cycle-promoting human Forkhead transcription factors	207
Palombo, Marta Investigation of NRF2 role in osteosarcoma	208
Panfilov, Mikhail Mechanistic insights into the sliding of individual nucleosomes under single-turnover conditions	209
Parassiadis, Christina Elucidating the mechanistic role of EZH2 in controlling pancreatic cancer cell identity	210
Park, Si Hoon Structural insight of crosstalk between transcription factors and Mediator complex	211
Patsis, Panos Dissecting the regulatory mechanisms of the histone H3 lysine 9 methyltransferase Clr4	212
Perez, Consuelo Genome-wide and multi-scale map of DNA supercoiling dynamics in differentiating human stem cells	213

Pirogov, Sergei Catalytic and non-catalytic functions of p300/CBP in zygotic genome activation	214
Pola, Victor Presenter: Montecino, Martin A. Epigenetic regulation supports dynamic changes in gene expression during early fear memory formation	215
Pollex, Tim Regulation of activity-dependent gene expression in neurons by CTCF and cohesin	216
Pouikli, Andromachi Mitochondrial dysregulation impacts transcription and splicing fidelity through epigenetic rewiring	217
Pownall, Mark Chromatin expansion microscopy reveals nanoscale organization of transcription and chromatin	218
Procaccia, Simone Only time will tell: observing regulatory dynamics at ultra-high temporal resolution with deep generative neural networks	219
Pudelko, Lorenz Investigating the role of chromatin organization in metabolic stress memory in C. elegans	220
Qureshi, Nusrat Presenter: Duss, Olivier Tracking transcription-translation coupling in real-time	221
Ramirez, Jose Miguel Cigarette smoking drives accelerated aging at the transcriptomic and epigenetic level	222
Rani, Harsha Enhancer reprogramming in p53 deficient colorectal cancer cells promotes cancer stemness and metastasis	223
Ravanelli, Giorgio Kinetic duality in transcription factors: a role in timing and levels decoupled regulation	224

Rekaik, Hocine Genetic ablation of Hox function in m reveals major rewiring in the early de		225
Reyes, Alejandro Degradation of WIZ reactivates fetal I globally accessible treatment for sick		226
Reynolds, Nicola Control of signal-responsive Enhanc remodelling proteins	Presenter: Hendrich, Brian er Resetting by chromatin	227
Rezende Pabst, Fernanda Investigating the role of metabolic en	nzymes in heterochromatin	228
Rogne, Marie FOXA1 SUMOylation affects its pione	Presenter: Eskeland, Ragnhild eering activity	229
Rosenkranz, Isabell Cold acclimation – a tug of war between and repression	een transcriptional activation	230
Rosok, Raya Interplay between deltaNp63 and GAS subtype identity in pancreatic cancer		231
Ross, Fiona Transcription dynamics underlying d mouse adult neural crest-derived nas		232
Rosti, Valentina Primary prostate cancer subtypes ide compartments profiling at time of dia		233
Salzler, Harmony R. Unique and redundant functions of c		234

Santulli, Martina High MYCN dosage reconfigures KAT2A/E2F3a intracellular interaction to foster Neuroblastoma cells towards more aggressive behaviors	235
Schaaf, Alina Chromatin and condensates in transcription regulation	236
Schwaemmle, Hanna Presenter: Braun, Simon CRISPR screen decodes SWI/SNF chromatin remodeling complex assembly	237
Schwager (Karpukhina), Anna A novel super-enhancer on derivative chromosome 14 contributes to mantle cell lymphoma pathogenesis through interchromosomal contacts	238
Seibert, Markus Presenter: Mermoud, Jacqueline Chromatin remodelling in fingerprint formation	239
Seman, Melissa Uncoupling the distinct functions of HP1 proteins during heterochromatin establishment and maintenance	240
Shah, Syed Zawar Structural characterisation of human RNA Polymerase III Pre-Initiation Complex (PIC)	241
Sharma, Shreya Investigating epigenetic alterations during epithelial to mesenchymal transition (EMT) in colorectal cancer cells with diverse p53 status	242
Sheikh, Nizamuddin Dissecting the mechanism of the BTAF1 and NC2 complex in disassembly of pol II pre-initiation complexes	243
Shlamovitz, Ivan Ubiquitylation of RNA Polymerase II controls early stages of the transcription cycle	244

Sidorov, Sviatoslav Cross-tissue isoform switches by human transcription factors suggest a widespread regulatory mechanism	245
Sloutskin, Anna Presenter: Juven-Gershon, Tamar From promoter motif to cardiac function: A single DPE motif affects transcription regulation and organ function in vivo	246
Snyder, Shannon Gene expression investigation of genetic and non-genetic variation of lifespan in Daphnia	247
Souto, Yara Bromodomain-specific Functions in Regulating Chromatin Interactions in PDAC	248
Steinhauser, Sebastian The transcription factor ZNF469 regulates collagen production in liver fibrosis	249
Stepien, Agata Pol II transcription termination is accelerated in colorectal cancer	250
Stocks, Jon Presenter: Gilbert, Nick Gene activation is abrogated by disrupting a SAF-A/RNA nuclear mesh	251
Stoos, Lisa Distinct mechanisms of chromatin engagement by transcriptional activators and repressors: insights from bHLH transcription factors	252
Stoy, Henriette Towards a genome-wide, uranyl-based DNA structure probing approach	253
Sudarshan, Nikita The interplay of Asr1 and Ser7 phosphorylation of RNA polymerase II CTD mediates sub telomeric gene silencing	254
Tan, Winnie	255

Poster	s A-Z
Tariq, Ammarah Structure and regulation of the H2B E3 ligase complex	256
Thomas, Henry Fabian Single-molecule analysis of HOX pioneer transcription factors	257
Toth, Daniel Marton Presenter: Aranyi, Tamas Hepatocyte-specific Dnmt3a and Dnmt3b knockout mice develop pre-cancerous phenotype	258
Traynor, Sarah A universal spike-in normalization strategy for CUT&RUN, CUT&Tag, and ATAC-Seq	259
Trendel, Jakob Mapping the human proteome with physical access to DNA	260
Trouth, Abby Presenter: Ramachandran, Srinivas G1 length dictates heterochromatin landscape	3 261
Trummel, Deborah Targeted protein depletion to investigate Drosophila u-shaped, a regulator of hematopoiesis	262
Umehara, Takashi How p300/CBP propagates histone acetylation	263
Unfried, Juan Pablo LncRNA FOSIL regulates FOS induction dynamics	264
Valakos, Dimitrios Dynamic reorganization of the macroH2A chromatin landscape during cellular reprogramming rewires gene expression programs towards pluripotency	265

266

Transcriptional regulation in mosquito immunity: insights from precision run-on sequencing (PRO-seq)

van Hout, Femke

van Zundert, Brigitte AJ Altered epigenetic landscape at the go methyltransferases Suv39h1/2 in ALS epi-editors to restore their expression	astrocytes, and engineering of	267
Vazquez Bolado, Alicia Uxia Transcriptional regulation in quiescen	nce cells	268
Vecchia, Luca Transcription factors OCT4-SOX2 regreemodeling directionality and activity	Presenter: Weiss, Joscha ulate SWI/SNF chromatin	269
Vizjak, Petra ISWI catalyzes nucleosome sliding in	Presenter: Mueller-Planitz, Felix condensed nucleosome arrays	270
Voichek, Yoav Widespread position-dependent enha	ncers in plants	271
Wiechens, Elina Backwards ahead: cooperating conve	Presenter: Fischer, Martin rgent promoters	272
Wiechens, Nicola Heterogeneity in the response to BAF	Presenter: Owen-Hughes, Tom complex inactivation	273
Wilson, Claire Investigating the role of CpG islands a in cancer	and DNA methylation at 3'UTRs	274
Xiao, Tianyi Oriented nucleosome binding by the μ	pioneer factor ELF2	275
Yanagisawa, Tatsuo Structural basis of eukaryotic transcri exonuclease complex	iption termination by the Rat1	276
Cancelled		277

Yue, Xue Global reorganization of chromatin fiber during stem cell differentiation	278
Zaratiegui, Carole Uncovering the role of mitochondrial stress in regulating the spatial organization of the genome	al 279
Zheng, Xin Presenter: Yin, Yimeng Impact of cytosine methylation on the binding of human transcript factors to nucleosome	ion 280
Zirden, Linda Proximity based interactomes of HSF1 under stress conditions	281
Zumer, Kristina FACT maintains chromatin architecture and thereby stimulates RN polymerase II pausing during transcription in vivo	A 282