

INTERNATIONAL MINI - CONFERENCE

CHROMOSOMES AND MITOSIS

December 8, 2017 Conference hall Institute of Molecular and Cellular Biology Lavrentiev Ave., 8/2 Novosibirsk, Russia

8:30 - 9:00	Conference participants registration
	Chairman: Gunter Reuter
9:00 – 10:00	Maurizio Gatti Department of Biology and Biotechnology 'C. Darwin', Sapienza, University of Rome, Rome, Italy; Institute of Molecular and Cellular Biology SB RAS, Novosibirsk, Russia The role of pendolino (peo) in the epigenetic regulation of Drosophila telomeres
10:00 – 10:30	Dmitry Zharkov Institute of Chemical Biology and Fundamental Medicine SB RAS, Novosibirsk, Russia; Novosibirsk State University, Novosibirsk, Russia DNA damage in uncharted waters of M phase
10:30 – 10:50	Anton Strunov Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia; Institute of Molecular and Cellular Biology SB RAS, Novosibirsk, Russia Endoplasmic reticulum and mitochondria behavior during Drosophila S2 cell mitosis
10:50 – 11:10	Coffee break
	Chairman: Jop Kind
11:10 – 11:55	Gunter Reuter Developmental Genetics, Institute of Biology, Martin Luther University Halle-Wittenberg, Halle, Germany The Su(var) gene complement and new chromatin functions in Drosophila
11:55 – 12:15	Stepan Belyakin Institute of Molecular and Cellular Biology SB RAS, Novosibirsk, Russia; Novosibirsk State University, Novosibirsk, Russia Functional dissection of Drosophila melanogaster SUUR protein influence on H3K27me3 profile
12:15 – 12:35	Evgeniya Andreyeva Institute of Molecular and Cellular Biology SB RAS, Novosibirsk, Russia Functional characterization of the conservative protein CG17337 in Drosophila melanogaster
12:35 – 12:55	Arcady Putilov Research Group for Math-Modeling of Biomedical Systems, the Research Institute for Molecular Biology and Biophysics, Novosibirsk, Russia How our clocks are evolving? Latitude-driven polygenic adaptation in favor of evening preference might occur by small allele frequency shifts spread across numerous loci in different circadian genes on different chromosomes
12:55 – 13:55	Lunch (not provided)
	Chairman: Maurizio Gatti
13:55 – 14:40	Jop Kind Hubrecht Institute, Koninklijke Nederlandse Akademie van Wetenschappen (KNAW), Utrecht, The Netherlands Untangling spatial genome organization: detailed insights from single-cell DamID
14:40 – 15:00	Alexey Pindyurin Institute of Molecular and Cellular Biology SB RAS, Novosibirsk, Russia Large fraction of heterochromatin in Drosophila neurons is simultaneously bound by B-type lamin and HP1a
15:00 – 15:30	Igor Sharakhov ^{Virginia Polytechnic Institute and State University, Blacksburg, VA, USA The interplay between chromosome-nuclear envelope attachments and chromosome-chromosome contacts}

15:30 – 15:50	Veniamin Fishman
	Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia; Novosibirsk State University, Novosibirsk, Russia
	Evolutionary analysis of 3-dimentional genome architecture

	Chairman: Igor Sharakhov
16:05 – 16:35	Irina Kolesnikova Institute of Molecular and Cellular Biology SB RAS, Novosibirsk, Russia Alteration of rRNA expression levels in patients with intellectual disability and enlarged p-arms on acrocentric autosomes
16:35 – 16:55	Stanislav Vasilyev Research Institute of Medical Genetics, Tomsk National Research Medical Center RAS, Tomsk, Russia DNA methylation in inherited copy number variations with incomplete penetrance
16:55 – 17:15	Renata Savchenko Research Institute of Medical Genetics, Tomsk National Research Medical Center RAS, Tomsk, Russia The role of the ADAMTS1, THBS1, WHSC1, RBFOX2 genes in the radiation-induced human somatic cellular response to DNA damage
17:15 – 17:35	Nikolay Skryabin Research Institute of Medical Genetics, Tomsk National Research Medical Center RAS, Tomsk, Russia Identification of loss of heterozygosity in miscarriages from families with recurrent pregnancy loss

Only the presenting authors are indicated; all co-authors of the presentations are listed in the abstract book.