

June 25 (Wednesday)

Time	Small Hall	Time	Exhibition Centre	Time	Music hall
09:00 – 13:00	BGRS-2014. Systems computational biology <i>Chairpersons:</i> <i>Prof. L. Kaderali, University of Technology Dresden, Germany</i> <i>Dr. A. Ratushny, Institute for Systems Biology, Seattle, USA</i> <i>Prof. V.P. Golubyatnikov, Sobolev Institute of Mathematics SB RAS, Novosibirsk, Russia</i>	09:00 – 13:00	ISHG-2014. MICROSYMPOSIUM “TRANSLATIONAL MEDICINE” <i>Chairpersons:</i> <i>Prof. M. Moshkin, Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia</i> <i>Prof. Y Iwakura, Research Institute for Biomedical Sciences, Tokyo University of Science, Tokyo, Japan</i>	09:00 – 13:00	School SBB-2014. Section “Evolutionary bioinformatics”
9:00-9:45	L. Kaderali University of Technology Dresden, Germany Systems biology of host-pathogen interactions KEYNOTE TALK	09:00-09:10	M.P. Moshkin Opening remark	9:00-9:45	Recommended to attend: L. Kaderali University of Technology Dresden, Germany Systems biology of host-pathogen interactions (Small Hall)
9:45-10:10	M. Djordjevic, K. Severinov, and M. Djordjevic Institute of Physics, University of Belgrade, Belgrade, Serbia Modeling bacterial immune systems: CRISPR/CAS regulation	10:00-10:20	M.Y. Pakharukova, N.I. Ershov, D.S. Pirozhkova, V.A. Vavilin, K.S. Zadesenets, T.G. Duzhak, V.A. Mordvinov Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Molecular mechanisms of metabolism, excretion and drug tolerance in human liver fluke <i>Opisthorchis felineus</i>	9:50-10:30	C. Mitra University of Hyderabad, Hyderabad, India Monte Carlo methods for metabolic kinetics
10:10-10:35	A. Barlukova, S. Honoré, F. Hubert, M. Petit Aix-Marseille University, France Dynamic instabilities of microtubules	10:20-10:40	E. Kozhevnikova Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Gut mucosal proteoglycan Mucin2 compromises animal sexual behavior		
10:35-11:00	M.A. Duk, A.M. Samsonov, M.G. Samsonova A.F. Ioffe Physical-technical Institute RAS, St.Petersburg, Russia Mechanism of miRNA action defines the dynamical behavior of miRNA-mediated feed-forward loops	10:40-11:00	P.N. Menshanov, A.E. Akulov, N.N. Dygalo Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Hippocampal neurochemical profile in neonatal rats: effects of anesthesia		

11:00-11:20 Coffee break					
11:20-12:05	M. Binder German Cancer Research Center, Division Virus-Associated Carcinogenesis, Heidelberg, Germany Modelling of the hepatitis C virus life cycle KEYNOTE TALK	11:20-11:40	S.P. Medvedev Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia The use of new genome engineering technologies for creation of isogenic amyotrophic lateral sclerosis cell model	11:20-12:05	Recommended to attend: M. Binder German Cancer Research Center, Heidelberg, Germany Modelling of the hepatitis C virus life cycle KEYNOTE TALK
12:05-12:30	N. Ivanisenko , E. Mishchenko, I. Akberdin, P. Demenkov, K. Kozlov, D. Todorov, V.V. Gursky, M.G. Samsonova, A.M. Samsonov, D. Clausznitzer, L. Kaderali, N.A. Kolchanov, V.A. Ivanisenko Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Mathematical model for subgenomic hepatitis C virus replication: impact of drug resistance	11:40-12:00	A.A. Nemudryi Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Rescue of Brattleboro rat strain mutant phenotype using TALEN and CRISPR/Cas9 systems		
12:30-12:45	N. Egorov InterLabService Advanced achievements of Illumina next- generation sequencing	12:00-12:20	O.S. Kozhevnikova , E.E. Korbolina, N.I. Ershov, N.G. Kolosova Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Aging of rat retina: transcriptome study	12:10-13:00	
		12:20-12:40	A.B. Salmina , N.A. Malinovskaya, O.L. Lopatina, Y.K. Khomleva Krasnoyarsk State Medical University, Krasnoyarsk, Russia Molecular markers and targets for diagnostics and treatment of neurodegeneration and neurodevelopmental disorders		D. Liberles Department of Molecular Biology, University of Wyoming, Laramie, USA The lecture on sequence analysis pipelines to uncover lineage- specific evolution.
13.00-14.00 Lunch					
14:00-17:15	BGRS-2014. Systems computational biology <i>Chairpersons:</i> <i>Prof. L. Kaderali, University of Technology Dresden, Germany</i> <i>Dr. A. Ratushny, Institute for Systems Biology, Seattle, USA</i> <i>Prof. V.P. Golubyatnikov, Institute of Mathematics SB RAS, Novosibirsk, Russia</i>	14:00-17:15	BGRS-2014. Plant systems biology <i>Chairperson:</i> <i>Prof. E. Salina, Institute of Cytology and Genetics SB RAS</i>	14:00-17:20	ISHG-2014. MICROSYMPOSIUM "TRANSLATIONAL MEDICINE" <i>Chairpersons:</i> <i>Prof. M. Moshkin, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i> <i>Prof. S. A. Brown, Institute of Pharmacology and Toxicology, University of Zurich, Switzerland</i>

14:00-14:25	K.N. Kozlov , V.V. Gursky, I.V. Kulakovskiy, V.V. Muzhichenko, M.G. Samsonova St.Petersburg State Polytechnical University, St.Petersburg, Russia Sequence -based model of gap gene regulatory network	14.00-14.35	M.V. Kapralov , S.M. Whitney Plant Science Division, Research School of Biology, The Australian National University, Canberra, Australia Evolution of Rubisco encoding genes in plants and its implications for Rubisco engineering in crops KEYNOTE TALK	14:00-14:40	S.A. Brown Institute of Pharmacology and Toxicology, University of Zurich, Zurich, Switzerland Mechanisms of circadian plasticity KEYNOTE TALK
14:25-14:50	E.V. Kashina , D.Y. Oshchepkov, E.A. Oshchepkova, A.G. Shilov, E.V. Antontseva, D.P. Furman, V.A. Mordvinov Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Dioxin-mediated regulation of genes involved in cytokines production by macrophages	14:35-14:55	P.M. Ponomarenko, M.P. Ponomarenko Children's Hospital Los Angeles, Los Angeles, USA An empirical equilibrium equation of a gene response to auxin in plants allows to predict quantitatively the auxin response	14:40-15:00	A. Baranova, L.Wang, T. Cui, B. Veytsman , S. Bruskin School of Systems Biology, George Mason University, Fairfax, USA Attractor based classifiers for prediction of post-treatment survival in cancer and detection of non-malignant diseases
14:50-15:15	I.R. Akberdin , T.V. Ermak, F.V. Kazantsev, T.M. Khlebodarova, V.A. Likhoshvai Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Kinetic modeling of pyrimidine biosynthesis is a first step to <i>in silico</i> bacterial cell	14:55-15:15	F.V. Kazantsev, V.V. Chernova, A.V. Doroshkov, N.A. Omelyanchuk, V.V. Mironova , V.A. Likhoshvai Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Self-organization mechanisms for auxin distribution in the root apical meristem	15:00-15:20	O.E. Redina , S.E. Smolenskaya, T.O. Abramova, L.N. Ivanova, A.L. Markel Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia Genetic dissection of inherited hypertensive state in ISIAH rats using kidney gene-expression and genome mapping
15:15-15:40	J. Narula, A. Kuchina, G.M. Süel and O.A. Igoshin Department of Bioengineering, Rice University, Houston, USA Slowdown of cell growth acts as the signal triggering cellular differentiation	15:15-15:35	A.I. Perfilyeva Siberian Institute of Plant Physiology and Biochemistry SB RAS, Irkutsk The statistical analysis of level of an expression of a series of proteins of a plant of <i>Arabidopsis thaliana</i> in the conditions of stressful influence	15:20-15:40	E. Pozhilenkova Krasnoyarsk State Medical University, Krasnoyarsk, Russia Development of cellular models for translational research in neuropharmacology
15:40-16:00 Coffee break					
16:00-16:25	H. Binder , H. Wirth, A. Arakelyan, K. Lembcke, E.S. Tiys, V. Ivanishenko, N.A. Kolchanov, A. Kononikhin, I. Popov, E.N. Nikolaev, L. Pastushkova, I.M. Larina Interdisciplinary Centre for Bioinformatics, Universität Leipzig, Leipzig, Germany A machine learning analysis of urine proteomics in space-flight simulations	16:00-16:20	I.V. Pinsky , A.T. Ivashchenko Al-Farabi Kazakh National University, Almaty, Kazakhstan miRNAS binding to mRNAs of rice Myb genes	16:00-16:20	I.L. Erokhin National Biotechnological Company, LLC, Moscow, Russia Oncogenesis model based on the genome structure of multicellular organisms

16:25-16:50	A. Kursanov , O. Solovyova, L. Katsnelson, K. Medvedev, A. Vasilyeva, N. Vikulova, V.S. Markhasin Institute of Immunology and Physiology UB of RAS, Yekaterinburg, Russia Cardiac mechanics, calcium overload and arrhythmogenesis	16:20-16:45	Y. Kanayama , H. Ikeda School of Agricultural Science, Tohoku University, Sendai, Japan Metabolome and transcriptome analyses of a tomato introgression line containing a <i>Solanum pennellii</i> chromosome segment	16:20-16:40	B. Veytsman , T.Cui, L. Wang, A. Baranova School of Systems Biology, George Mason University, Fairfax, USA Systemic shifts in micro RNA landscape as a diagnostic and prediction tool
16:50-17:15	P.A. Ryzhkov , N.S. Ryzhkova Scientific Research Institute of Biology SFEDU, Rostov-on-Don, Russia Graph model of type I diabetes	16:45-17:05	D.A. Afonnikov , M.A. Genaev, E.G. Komyshev, A.V. Doroshkov, T.A. Pshenichnikova, E.V. Morozova, A.V. Simonov ICG SB RAS, Novosibirsk, Russia Computer high-throughput approaches to wheat phenotyping	16:40-17:00	L.N. Grinkevich I.P. Pavlov Institute of Physiology of RAS, St. Petersburg, Russia Epigenetic mechanisms of memory formation: the role of the histone acetylation and methylation in aversive learning
17:15-17:50	S.A. Lashin , Yu.G. Matushkin Institute of Cytology and Genetics SB RAS; Novosibirsk State University, Novosibirsk, Russia DEC: software tools for simulation evolution in diploid populations	17:05-17:25	A.V. Doroshkov , U.S. Zubairova, M.A. Genaev, S.V. Nikolaev, T.A. Pshenichnikova, D.A. Afonnikov ICG SB RAS, Novosibirsk, Russia Analysis of bread wheat leaf pubescence formation and diversity using image analysis technique and mathematical modeling	17:00-18:00	School BB-2014. Practical training (ICG SB RAS: Lavrentyeva, 10) F.V. Kazantsev Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia
17:50-18:10	O.V. Popik , N.A. Kolchanov, V.A. Ivanisenko Center of neurobiology and neurogenetics, ICG SB RAS, Novosibirsk, Russia Mathematical modeling of the interactions between molecular genetic systems	17:25-17:45	K.N. Sarsenbayev Eurasian National University, Astana, Kazakhstan Proteomic information of spring wheat varieties differing in resistance to infection after <i>Puccinia recondita</i> inoculation		
		17:45-18:10	E.M. Sergeeva , E.M. Timonova, L.L. Bildanova, M.K. Koltunova, M.A. Nesterov, F. Magni, Z. Frenkel, J. Dolezel, J. Faris, P. Sourdille, C. Feuillet, E.A. Salina ICG SB RAS, Novosibirsk, Russia The progress in physical mapping of chromosome 5b of bread wheat <i>Triticum aestivum</i>		
18:10-19:00	Foyer of Small Hall (House of Scientists SB RAS) POSTER SESSION				