

## June 27 (Friday)

e	Small Hall	Time	Exhibition Centre	Time	Music hall
09:00 – 12:35	<b>BGRS-2014. Proteomics, metabolomics and computational pharmacology</b> <i>Chairpersons:</i> <i>Prof. E. Nikolaev, Institute for Biomedical Problems of RAS, Moscow, Russia</i> <i>Prof. I. Larina, Institute for Biomedical Problems of RAS, Moscow, Russia</i>	09:00 – 12:15	<b>MM-HPC-2014. Pharmacokinetics and immunology</b> <i>Chairpersons:</i> <i>Prof. A.I. Ilyin, O.I. Krivorotko</i> <i>Institute of Computational Mathematics and Mathematical Geophysics SB RAS, Novosibirsk, Russia</i>	09:00 – 13:00	<b>School SBB-2014.</b> <b>Presentations by young scientists</b>
09:00-09:45	<b>I.M. Larina, V.A. Ivanisenko, E.N. Nikolaev</b> Institute for Biomedical Problems of RAS, Moscow, Russia <b>Proteomics extreme impacts as a tool for systems biology</b> <b>KEYNOTE TALK</b>	09:00-09:30	<b>A.I. Ilyin, S.I. Kabanikhin, D.A. Voronov</b> Institute of Computational Mathematics and Mathematical Geophysics SB RAS <b>Numerical solutions of inverse problem of pharmacokinetics. Identifiability of compartmental models</b> <b>KEYNOTE TALK</b>	09:00-09:20	<b>V.V. Lavrekha, N.A. Omelyanchuk, V.V. Mironova</b> Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia <b>Modeling of cell dynamics in the root apical meristem with dynamical grammar</b>
		09:30-09:50	<b>A.I. Ilyin, S.I. Kabanikhin, O.I. Krivorotko</b> Institute of Computational Mathematics and Mathematical Geophysics SB RAS <b>The identification and refinement of parameters of mathematical models in immunology</b>	09:20-09:40	<b>S. Maltseva, A.P. Chupakhin, A.A. Cherevko, A.K. Khe, E.U. Derevtsov, A.E. Akulov</b> Institute of Mathematics SB RAS, Novosibirsk, Russia <b>Reconstrucion of the mouse brain vasculature according to the data of high-field MRI scanner</b>
09:45-10:10	<b>T. Duzhak, Yu.P. Tsentalovich</b> International Tomography Center SB RAS, Novosibirsk, Russia <b>Lens cataract: effects of crystallins modification</b>	09:50-10:10	<b>M.A. Shishlenin</b> Sobolev Institute of Mathematics SB RAS, Novosibirsk, Russia <b>Continuation of the acoustic field in tomography</b>	09:40-10:00	<b>N. Nikitina, E.E. Ivashko, Y. Gupta, R. Lüdwig, S. Möller.</b> Institute of Applied Mathematical Research Karelian Research Center RAS, Petrozavodsk, Russia <b>Boinc-based desktop GRID infrastructure for virtual drug screening</b>
10:10-10:35	<b>M.A. Karpova, D.A. Karpov, M.V. Ivanov, V.G. Zgoda, M.V. Gorshkov, S.A. Moshkovskii</b> Orekhovich Institute of Biomedical Chemistry RAMS, Moscow, Russia <b>Cancer cell line recognition by shotgun</b>	10:10-10:30	<b>A.A. Vitvitsky</b> ICM&MG SB RAS, Novosibirsk, Russia <b>Computer simulation of self-organization in the bacterial MINCDE system</b>	10:00-10:20	<b>V. Mironova, D. Novikova</b> Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia <b>Auxin-induced changes of transcriptome in <i>Arabidopsis thaliana</i> l. roots.</b>

	<b>proteomics using cancer exome data</b>	10:30-10:50	<b>A.A. Polshchitsin</b> , V.M. Nekrasov, A.V. Chernyshev, V.P Maltsev Institute of chemical kinetics and combustion; Novosibirsk State University, Novosibirsk, Russia. <b>Solution of inverse immunoagglutination kinetics problem for patchy particles with a small number of binding sites</b>	10:20-10:40	<b>I. Protsyuk</b> UniPro (Новосибирск), Россия <b>Shared bioinformatics databases within Unipro UGENE</b>
10:35-11:00	I.A. Mescheryakova, E.V. Demidova, E.A. Demidov, T.N. Goryachkovskaya, V.M. Popik, A.I. Semenov, G.N. Kulipanov, N.A. Kolchanov, <b>S.E. Peltek</b> Institute of Cytology and genetics SB RAS, Novosibirsk, Russia <b>Genomic proteomics profiling</b>			10:40-11:00	<b>O. Perfilyeva</b> ICG SB RAS, Novosibirsk, Russia <b>Mathematical modeling of peptidoglycan precursor biosynthesis in the cytoplasm of Esherichia coli cell</b>
11:00-11:20 <b>Coffee break</b>					
11:20-11:45	<b>L.H. Pastushkova</b> , A.S. Kononihin, E.S. Tiys, O.A. Obraztsova, I.V. Dobrohotov, K.S. Kireev, V.A. Ivanisenko, E.N. Nikolaev, I. M. Larina Russian Federation State Scientific Center - Institute for Biomedical Problems of RAS; Moscow, Russia <b>Identifying overrepresented biological processes in cosmonauts on the first day</b>	11:15-11:35	<b>A. Levichev</b> , A. Palyanov Sobolev Institute of Mathematics SB RAS, Novosibirsk, Russia <b>On a modification of the theoretical basis of the Penrose-Hameroff model of consciousness</b>	11:20-11:40	<b>I. Pinskyi</b> , A. Ivashchenko Al-Farabi Kazakh National University, Almaty, Kazakhstan <b>miRNAs binding to mRNAs of rice Myb genes</b>
		11:35-11:55	<b>A. Yu. Pyrkova</b> , A.T. Ivashchenko, O.A. Berillo Al-Farabi Kazakh National University, Almaty, Kazakhstan <b>Modelling of the problem of multiple alignment of the nucleotide sequences and dendrogram construction</b>	11:40-12:00	<b>N.V. Sviridova</b> , K. Sakai Computing Center FEB RAS, Khabarovsk, Russia <b>Application of nonlinear time series analysis for hemodynamic model validation on the base of photoplethysmogram signal</b>
11:45-12:10	<b>P.S. Sherin</b> , E.A. Zelentsova, E.D. Sormacheva, T.G. Duzhak, Yu.P. Tsentalovich International Tomography Center SB RAS, Novosibirsk, Russia <b>UVA-induced modifications of lens Alpha-crystallin</b>	11:55-12:15	<b>A.A. Bedelbayev</b> Institute of Mathematics and Mechanics KazNU, Almaty, Kazakhstan <b>Computer mathematical and biochemical modeling and simulation of the life processes in human organs</b>	12:00-12:20	<b>N. Safronova</b> , Y.L. Orlov ICG SB RAS, Novosibirsk, Russia <b>Computer analysis of human SNP containing sites by methods of text complexity estimations</b>
				12:20-12:40	<b>A. Sergeev</b> Institute of Mathematical Problems of Biology RAS, Puschino, Russia <b>Graph database for molecular biology- advantages of the graph representation of data</b>
13:00-14:00 <b>Lunch</b>					

14:00-17:50	<b>BGRS-2014. Proteomics, metabolomics and computational pharmacology</b> <i>Chairpersons:</i> <i>Prof. E. Nikolaev, Institute for Biomedical Problems of RAS, Moscow, Russia</i> <i>Prof. I. Larina, Institute for Biomedical Problems of RAS, Moscow, Russia</i>	14:00-17:55	<b>MM-HPC-2014. Gene networks</b> <i>Chairpersons:</i> <i>Prof. V.P. Golubyatnikov, Dr. N.B. Ayupova, Institute of Computational Mathematics and Mathematical Geophysics SB RAS, Novosibirsk, Russia</i>	14:00-17:00	<b>School SBB-2014.</b>  <b>Presentations by young scientists.</b>  <b>Nomination of best presentations</b>
14:00-14:25	<b>S.F. Chekmarev</b> Institute of Themophysics SB RAS; Novosibirsk State University, Novosibirsk, Russia <b>Driving force for protein folding: the two-component potential</b>	14:00-14:30	N.B.Ayupova, <b>V.P. Golubyatnikov</b> Sobolev Institute of Mathematics SB RAS; Novosibirsk State University, Novosibirsk, Russia <b>On geometry of phase portraits of some low-dimensional gene network models</b> <b>KEYNOTE TALK</b>	14:00-14:20	<b>V. Sokolov, U.G. Matushkin</b> ICG SB RAS, Novosibirsk, Russia <b>Analysis of bacteria and archaea genomes available in GenBank database by “Eloe” program</b>
14:25-14:50	<b>A.O. Chugunov, P.E. Volynsky, R.G. Efremov</b> Shemyakin-Ovchinnikov Institute of Bio-organic Chemistry RAS, Moscow, Russia <b>High-performance computing provides insight into the innermost organization of procaryotic membranes</b>	14:30-14:50	<b>F.V. Kazantsev, I.R. Akberdin, N.L. Podkolodnyy, V.A. Likhoshvai</b> Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia <b>Gene networks modeling: specification language</b>	14:20-14:40	<b>A. Spitsina, V.M. Efimov, V.N. Babenko, Y.L. Orlov</b> ICG SB RAS, Novosibirsk, Russia <b>Computer analysis of human gene expression data using BioGPS database of Affymetrix microarrays</b>
				14:40-15:00	<b>T.S. Troeglazova, D.Ja. Baishibaev, A.V. Penenko, S.V. Nikolaev, U. Zubairova</b> Novosibirsk State University, Russia <b>On a parallel algorithm for morpho-gene diffusion-reaction processes simulation on a 2D cell ensemble</b>
14:50-15:15	<b>A.Y. Nyporko</b> Taras Shevchenko National University of Kyiv, Kyiv, Ukraine <b>Features 8-oxo-DGTP behavior in active site of human DNA polymerase</b>	14:50-15:10	<b>S.V. Lenskiy, T.I. Lenskaya</b> Ural Federal University, Ekaterinburg, Russia <b>Power law for rank distribution of gene density in human genome projects</b>	15:00-15:20	<b>V. Chernova, A. Doroshkov, V. Mironova, N. Omelyanchuk</b> ICG SB RAS, Novosibirsk, Russia <b>PIN-transporters in the root meristem of Arabidopsis thaliana I- image analysis of expression patterns</b>
15:15-15:40	<b>K.E. Medvedev, D.A. Afonnikov</b> Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia <b>Computational investigation of high pressure and temperature influence</b>	15:10-15:30	A.A. Akinshin, T.A. Bukharina, D.P. Furman, <b>V.P. Golubyatnikov</b> Sobolev Institute of Mathematics SB RAS, Novosibirsk, Russia <b>Modeling of two cells complex in morphogenesis of D. Melanogaste mechanoreceptors</b>		

15:40-16:05	<b>E.A. Aksianov</b> , A.V. Alexeevsky Belozersky Institute of Physico-Chemical Biology, MSU, Moscow, Russia <b>Alignment of “unalignable” protein structures</b>	15:30-15:50	<b>V.Yu. Salamatova</b> Moscow Institute of Physics and Technology, Moscow, Russia <b>Modelling of soft tissues deformation. Alternative approaches.</b>	15:40-16:00	AWARD FOR THE BEST YOUNG SCIENTISTS REPORTS
16:05-16:25 <b>Coffee Break</b>					
16:25-16:50	<b>A.A. Anashkina</b> , N.G. Esipova, E.N. Kuznetsov, V.G. Tumanyan Engelhardt Institute of Molecular Biology RASs, Moscow, Russia <b>Native proteins and decoys: subtle structure differences</b>	16:15-16:35	<b>B. Karacaören</b> Department of Animal Science, Akdeniz University, Antalya, Turkey <b>Dynamic association mapping based on random walk model using simulated QTLMAS data set</b>		
16:50-17:15	A.M. Andrianov, I.A. Kashyn, <b>A.V. Tuzikov</b> United Institute of Informatics Problems NAS Belarus, Minsk, Belarus <b>Computational prediction of novel anti-HIV-1 agents based on potent and broad neutralizing antibody VRC01</b>	16:35-16:55	<b>L.A. Krasnobaeva</b> , L.V. Yakushevich Siberian State Medical University; Tomsk State University, Tomsk, Russia <b>Rotational dynamics of bases in the gene coding Interferon alpha 17 (IFNA17)</b>		
		16:55-17:15	<b>E.S. Fomin</b> ICG SB RAS, Novosibirsk, Russia <b>Reconstruction cyclic sequences from their circular distances multiset</b>		
		17:15-18:00	Auditorium near Conference hall in the Exhibition Centre SB RAS <b>POSTER SESSION</b>		
<b>18:30 - 20:30</b> The closing ceremony of the Multi-Conference. Venue: Technopark of Novosibirsk Akademgorodok (Nikolaeva str, 11, 13th floor). Departure to the ceremony will be from the House of Scientists (left wing) at 18:00. Entrance to the ceremony by badges. .....					
<b>18:30 – 20:30</b> Церемония закрытия Мульти-конференции. Технопарк Академгородка (ул. Николаева, 11, 13-ый этаж). Отъезд участников на церемонию – в 18:00 от Дома Ученых (левое крыло). Вход по бейджам.					