

# SYMPOSIUM

## Genomics, transcriptomics, bioinformatics

Oral reports

7 July, Tuesday

Big Conference Hall

Morning session 1. Genomics, transcriptomics, bioinformatics:  
Computational genomics and oncogenomics

Chair: Yakov Tsepilov, *Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

09:00 –  
09:30

### PLENARY REPORT

Understanding heterozygous carrying of deleterious variants as a key for personalized longevity

*Ancha Baranova,*  
School of Systems Biology, George Mason University, Fairfax, VA

10:25 –  
10:45

### Search of New Type of Spatial Organization of Nucleic Acids in Human Genome

Anastasia Zamoskovtseva<sup>1,2</sup>, Marsel Kabilov<sup>1</sup>, Alexander Lomzov<sup>1</sup>, Dmitrii Pyshnyi<sup>1</sup>

<sup>1</sup>*ICBFM SB RAS, Novosibirsk, Russia*

<sup>2</sup>*NSU, Novosibirsk, Russia*

10:45 –  
11:05

### Bioinformatic methods applied to the analysis of the genes retained after the whole genome duplication events in the sterlet genome (*Acipenser ruthenus*)

Mikhail Fofanov<sup>1</sup>, Tatyana Sheglova<sup>2</sup>, Vladimir Trifonov<sup>2</sup>, Manfred Schartl<sup>3</sup>

<sup>1</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>2</sup>*Institute of Molecular and Cellular Biology, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*University of Würzburg, Würzburg, Germany*

11:05 –  
11:25 **Coffee break**

11:25 – **GPU Based Composite Elements Discovery In Large DNA Datasets**  
11:45 Oleg Vishnevsky<sup>1,2</sup>, Andrey Bocharnikov<sup>2</sup>, Nikolay Kolchanov<sup>1,2</sup>  
<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

11:45 – **Using fast homology search tools for protein sequence functional annotation: a comparison**  
12:00 Pronozin Artem, Mikhail Genaev, Dmitry Afonnikov  
*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

12:00 – **Web – 3DPredictor: a Web Interface for High – resolution Prediction of Genome Architecture**  
12:20 Emil Valeev<sup>1</sup>, Polina Belokopytova<sup>1</sup>, Veniamin Fishman<sup>1,2</sup>  
<sup>1</sup>*Institute of Cytology and Genetics Novosibirsk, Russia*  
<sup>2</sup>*NSU, Novosibirsk, Russia*

**Lunch**

**Evening session 1. Genomics, transcriptomics, bioinformatics:  
Quantitative genetics and genomic epidemiology**

**Chairs: Georgii Bazykin, Skoltech, Moscow, Russia; Institute for Information Transmission Problems (Kharkevich Institute) of the Russian Academy of Sciences, Moscow, Russia;  
Vsevolod Makeev, Vavilov Institute of General Genetics, Moscow, Russia;  
Ivan Kylakovskiy, Engelhardt Institute of Molecular Biology RAS, Vavilov Institute of General Genetics RAS, Moscow, Russia;  
Yakov Tsepilov, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia.**

15:00 – 15:30 **Quantitative genetics and computational functional genomics as tools to study biology**  
Yurii Aulchenko  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
*Novosibirsk State University, Novosibirsk, Russia*

15:30 – 15:50 **Loci and genes involved in chronic musculoskeletal pain identified via analysis of genetically independent pain phenotypes**  
Yakov Tsepilov  
Yakov A. Tsepilov<sup>1</sup>, Sodbo Z. Sharapov<sup>1</sup>, Lennart C. Karssen<sup>2</sup>, Yurii S. Aulchenko<sup>3</sup>, Maxim B. Freidin<sup>4</sup>, Elizaveta E. Elgaeva<sup>1</sup>, Pradeep Suri<sup>5</sup>, Alexandra S. Shadrina<sup>1</sup>, Jan van Zundert<sup>6</sup>, Frances M.K. Williams<sup>4</sup>  
<sup>1</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>2</sup>PolyOmica, 's – Hertogenbosch, the Netherlands  
<sup>3</sup>Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia  
<sup>4</sup>King's College London, London, UK  
<sup>5</sup>VA Puget Sound Health Care System, Seattle, USA  
<sup>6</sup>Maastricht University Medical Centre, Maastricht, The Netherlands

15:50 – 16:05

**Labchip GX Touch Nucleic Acid Analyzer for quantitative assays and QC in Genomics**  
Ilse Villmann, Application Scientist, Perkin Elmer, United States  
**Sponsor report – BioLine**

16:05 – 16:30

**Quantitative genetics of protein N – glycosylation**  
Lucija Klaric<sup>1,2</sup>, Yurii S. Aulchenko<sup>3,4,5</sup>, Yakov A. Tsepilov<sup>3,5</sup>, Gordan Lauc<sup>2,6</sup>, Chloe M. Stanton<sup>1</sup>, Caroline Hayward<sup>1</sup>  
<sup>1</sup>University of Edinburgh, Edinburgh, United Kingdom  
<sup>2</sup>Genos Glycoscience Research Laboratory, Zagreb, Croatia  
<sup>3</sup>Institute of Cytology and Genetics of the SBRAS, Novosibirsk, Russia  
<sup>4</sup>PolyOmica, 's – Hertogenbosch, The Netherlands  
<sup>5</sup>Novosibirsk State University Novosibirsk, Russia  
<sup>6</sup>University of Zagreb, Zagreb, Croatia

16:30 – 16:45

**Results of genome – wide association study of plasma proteome N – glycosylation in 10,000 sample**  
Sodbo Sharapov<sup>1</sup>, Sofya Feoktistova<sup>1</sup>, Lucija Klaric<sup>2</sup>, Harry Campbell<sup>2</sup>, Matthias Schulze<sup>3</sup>, Yurii Aulchenko<sup>1</sup>, Yakov A. Tsepilov<sup>4</sup>, Eugene Tiys<sup>1</sup>, Karsten Suhre<sup>5</sup>, Malcolm Dunlop<sup>2</sup>, Tim Spector<sup>6</sup>, Elizaveta E. Elgaeva<sup>4</sup>, Frano Vuckovic<sup>7</sup>, Nishi Chaturvedi<sup>8</sup>, Frances Williams<sup>6</sup>, Gordan Lauc<sup>7</sup>  
<sup>1</sup>Institute of Cytology and Genetics Novosibirsk, Russia  
<sup>2</sup>University of Edinburgh, Edinburgh, United Kingdom  
<sup>3</sup>German Institute of Human Nutrition Potsdam – Rehbruecke. Nuthetal, Germany  
<sup>4</sup>Novosibirsk State University Novosibirsk, Russia  
<sup>5</sup>Weill Cornell Medicine – Qatar, Doha, Qatar  
<sup>6</sup>School of Life Course Sciences King's College London, London, United Kingdom  
<sup>7</sup>Genos Glycoscience Research Laboratory, Zagreb, Croatia  
<sup>8</sup>MRC Unit for Lifelong Hlth & Ageing University College London, London, United Kingdom

16:45 – 17:00

**A study of causal relationships between human IgG N – glycosylation traits and twelve associated diseases**  
Olga O. Zaytseva<sup>1</sup>, Gordan Lauc<sup>1</sup>, Sodbo Z. Sharapov<sup>2</sup>, Yakov A. Tsepilov<sup>3</sup>, Lucija Klaric<sup>4</sup>  
<sup>1</sup>Genos Glycoscience Research Laboratory, Zagreb, Croatia  
<sup>2</sup>Institute of Cytology and Genetics Novosibirsk, Russia  
<sup>3</sup>Novosibirsk State University Novosibirsk, Russia  
<sup>4</sup>University of Edinburgh, Edinburgh, United Kingdom

17:00 – 17:10

**Coffee break**

17:10 – 17:40

**Keynote report**

**Genomic epidemiology of SARS – CoV – 2 in Russia**

A. Komissarov<sup>1</sup>, K. Safina<sup>2</sup>, A. Fadeev<sup>1</sup>, D. Danilenko<sup>1</sup>, S. Garushyants<sup>3</sup>, V. Shchur<sup>4</sup>, G. Bazykin<sup>2,3</sup>

<sup>1</sup>*Smorodintsev Research Institute of Influenza*

<sup>2</sup>*Skolkovo Institute of Science and Technology (Skoltech)*

<sup>3</sup>*A.A. Kharkevitch Institute for Information Transmission Problems of the RAS*

<sup>4</sup>*Higher School of Economics National Research University*

17:40 – 17:55

**The role of host genetics in severity of COVID – 19**

Ivan Kuznetsov

Skolkovo Institute of Science and Technology, Moscow, Russia

*Novosibirsk State University, Novosibirsk, Russia*

17:55 – 18:05

**Short break**

18:20 – 18:35

**GWAS – MAP: the platform for analysis of results of genome – wide association studies**

Tatiana Shashkova<sup>1</sup>, Sodbo Sharapov<sup>1</sup>, Denis Gorev<sup>1</sup>, Yakov Tsepilov<sup>1</sup>, Yurii Aulchenko<sup>1</sup>, Eugene Pakhomov<sup>1</sup>, Lennart Karsen<sup>2</sup>

<sup>1</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>2</sup>*PolyKnomics's – Hertogenbosch, Netherlands*

## 8 July, Wednesday

Big Conference Hall

Morning session 2. Genomics and transcriptomics

Chair: Veniamin Fishman, *Institute of Cytology and Genetics Novosibirsk, Russia; NSU, Novosibirsk, Russia*

9.00 – 9.30

**PLENARY REPORT**

**Human Genome, Anesthesiology and Critical Care. 20 Years Later**

Vladimir Zelman,

*Keck School of Medicine - USC*

11:20 – 11:40

**Repetitive elements in the genome of Siberian larch (*Larix sibirica* Ledeb.)**

K.A. Miroshnikova<sup>1</sup>, M.G. Sadovsky<sup>3</sup>, V.S. Akulova<sup>3</sup>, V.V. Biriukov<sup>3</sup>, E.I. Bondar<sup>3</sup>, V.V. Sharov<sup>3</sup>, D.A. Kuzmin<sup>2</sup>, Y.A. Putintseva<sup>2</sup>, N.V.

Oreshkova<sup>3</sup>, K.V. Krutovsky<sup>2,4,5</sup>

<sup>1</sup>*Institute of Biophysics SB RAS, Krasnoyarsk, Russia*

<sup>2</sup> *SibFU, Krasnoyarsk, Russia*

<sup>3</sup> *FRC KSC SB RAS, Krasnoyarsk, Russia*

<sup>4</sup>Georg – August University of Göttingen, Göttingen, Germany

<sup>5</sup>Vavilov Institute of General Genetics, Moscow, Russia

11:40 – 12:00

**Analysis tandem repeats and retrotransposons of *Shepherdia argentea* (Pursh) Nutt**

Karina Bone<sup>1,2</sup>, Olga Razumova<sup>1,3</sup>, Gennady Karlov<sup>1</sup>, Ilya Kirov<sup>1</sup>

<sup>1</sup>All – Russia Research Institute of Agricultural Biotechnology, Moscow, Russia

<sup>2</sup>Russian State Agrarian University – Moscow Timiryazev Agricultural Academy, Moscow, Russia

<sup>3</sup>Kurchatov Genomic Center, Moscow, Russia

12:00 – 12:20

**Break**

12:20 – 12:35

**Analysis of the Complete Genome Sequence of Strain Concept – 8, the New Representative of the Genus *Methylococcus***

I.Y. Oshkin<sup>1,2</sup>, K.K. Miroshnikov<sup>1,2</sup>, D.V. Chernushkin<sup>3</sup>, N.V. Ravin<sup>2,4</sup>, V.O. Popov<sup>2</sup>, V.N. Khmelenina<sup>5,6</sup>, S.E. Belova<sup>1,2</sup>, A.V. Beletsky<sup>4,2</sup>, S.N. Dedysh<sup>1,2</sup>, S. Y. But<sup>5,6</sup>, N.S. Khokhlachev<sup>7</sup>, A.V. Mardanov<sup>2,4</sup>, N.V. Pimenov<sup>1,2</sup>

<sup>1</sup>Winogradsky Institute of Microbiology, Moscow, Russia

<sup>2</sup>Research Center of Biotechnology of the Russian Academy of Sciences, Moscow, Russia

<sup>3</sup>BIOSINTEZ, LLC

<sup>4</sup>Institute of Bioengineering, Moscow, Russia

<sup>5</sup>Federal Research Center "Pushchino Scientific Center for Biological Research of the Russian Academy of Sciences",

<sup>6</sup>G.K. Skryabin Institute of Biochemistry and Physiology of Microorganisms, Russian Academy of Sciences

<sup>7</sup>Gazprom VNIIGAZ

12:35 – 12:50

**Effective sample preparation for NGS – increasing productivity, reducing costs**

Baybaev Nikolay, *Dia – M*, Moscow, Russia

**Sponsor report – *Dia – M***

**Lunch**

**Evening session 2. Genomics, transcriptomics, bioinformatics:  
Gene regulation**

**Chairs: Yurii Aulchenko, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia; Novosibirsk State University, Novosibirsk, Russia  
Georgii Bazykin, Skoltech, Moscow, Russia; Institute for Information Transmission Problems (Kharkevich Institute) of the Russian Academy of Sciences, Moscow, Russia**

**Ivan Kylakovsky, Engelhardt Institute of Molecular Biology RAS, Vavilov Institute of General Genetics RAS, Moscow, Russia**

14.30 – 15.00

**PLENARY REPORT**

**Understanding the evolution of complex regions in bird genomes**

Alexander Sang-Jae Suh,  
*Department of Evolutionary Biology, Uppsala University, Uppsala, Sweden*

15:00 – 15:30

**Keynote report**

**Exploring the universe of transcription factor binding motifs in DNA**

Vsevolod Makeev

*Vavilov Institute of General Genetics, Moscow, Russia*

15:30 – 15:50

**Analysis of motifs co – occurrence in ChIP – seq data**

Victor Levitsky, Elena Zemlyanskaya, Dmitry Oshchepkov, Anton Tsukanov and Tatyana Merkulova

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

15:50 – 16:00

**Short break**

16:00 – 16:30

**AD ASTRA: the database of Allelic Dosage – corrected Allele – Specific TRAnscription factor binding suggests causal regulatory sequence variants of pathologies**

Sergey Abramov<sup>1</sup>, Eugene Baulin<sup>2</sup>, Vsevolod J Makeev<sup>1</sup>, Alexandr Boytsov<sup>1</sup>, Ivan Yevshin<sup>3</sup>, Ivan Kulakovsky<sup>4</sup>, Bykova Dariia<sup>5</sup>, Fedor Kolpakov<sup>6</sup>

<sup>1</sup>*Vavilov Institute of General Genetics RAS, Moscow, Russia*

<sup>2</sup>*Institute of Mathematical Problems of Biology RAS – the Branch of Keldysh Institute of Applied Mathematics of Russian Academy of Sciences, Pushchino, Russia*

<sup>3</sup>*BIOSOFT.RU LLC, Novosibirsk, Russia*

<sup>4</sup>*Engelhardt Institute of Molecular Biology RAS, Moscow, Russia*

<sup>5</sup>*Lomonosov Moscow State University, Moscow, Russia*

<sup>6</sup>*Institute of Computational Technologies SB RAS, Novosibirsk, Russia*

16:30 – 16:50

**Diversity of Cis – elements in Response to Dioxin in Human**

Evgenia Oshchepkova<sup>1</sup>, Yana Sizentsova<sup>1</sup>, Victoria Mironova<sup>1,2</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*NSU, Novosibirsk, Russia*

16:50 – 17:10

**Coffee break**

17:10 – 17:40

**GTRD – an integrated view on transcription regulation**

Fedor A. Kolpakov<sup>1,2</sup>, Ivan S. Evshin<sup>1,2</sup>, Semyon K. Kolmykov<sup>1,3</sup>, Yury V. Kondrakhin<sup>1</sup>, Mikhail A. Kulyashov<sup>1,4</sup>, Ruslan N. Sharipov<sup>1,2,4</sup>

<sup>1</sup>*Institute of Computational Technologies, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*BIOSOFT.RU, LLC, Novosibirsk, Russia*

<sup>3</sup>*FRC Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>4</sup>*Novosibirsk State University, Novosibirsk, Russia*

- 17:40 – 17:55**      **Meta – analysis of ChIP – seq Datasets Through Rank Aggregation Approach**  
Semyon K. Kolmykov<sup>1,2</sup>, Ivan S. Yevshi<sup>2,3</sup>, Yury V. Kondrakhin<sup>2</sup>, Anna S. Ryabova<sup>2,3</sup>, Ruslan N. Sharipov<sup>3</sup>, Fedor A. Kolpakov<sup>2,3</sup>  
<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
<sup>2</sup>*Institute of Computational Technologies SB RAS, Novosibirsk, Russia*  
<sup>3</sup>*BIOSOFT.RU, LLC, Novosibirsk, Russia*
- 17:55 – 18:10**      **News in microdissection in the context of transcriptomic studies and single cell research: AccuLift Fluidigm**  
Anna Tarasevich, *Helicon Company, Moscow, Russian Federation*  
**Sponsor report – Helicon Company**
- 18:10 – 18:25**      **JetGene – an Internet Resource for Analysis of Regulatory Regions or Nucleotide Contexts at Differently Translated Transcripts**  
N.S. Sadovskaya<sup>1</sup>, O.N. Mustafaev<sup>2</sup>, I.V. Goldenkova – Pavlova<sup>1</sup>, A.A. Tyurin<sup>1</sup>  
<sup>1</sup>*Tmiryazev Institute of Plant Physiology, RAS, Moscow, Russia*  
<sup>2</sup>*Genetic Resources Institute, ANAS, Baku, Azerbaijan*
- 18:25 – 18:40**      **New Approach to Genome – Wide Automated Inference of Bacterial Transcription Factor Binding Sites**  
Yevgeny Nikolaichik, Pavel Vychik  
*Belarusian State University, Minsk, Belarus*
- 18:40 – 18:55**      **Random Projections for functional signal extraction from single – cell RNA – seq data**  
Alexey Samosyuk  
*Skolkovo Institute of Science and Technology, Moscow, Russia*

## Poster session

### Statistical problems of clusters of transcription factor binding sites in plant genomes

Artur Dergilev<sup>1,2</sup>, Yuriy L. Orlov<sup>1,3</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*I.M.Sechenov First Moscow State Medical University, Moscow, Russia*

### Loci and genes involved in chronic musculoskeletal pain identified via analysis of genetically independent pain phenotypes

Yakov Tsepilov<sup>1</sup>, Maxim B. Freidin<sup>4</sup>, Alexandra S. Shadrina<sup>1</sup>, Sodbo Z. Sharapov<sup>1</sup>, Elizaveta E. Elgaeva<sup>1</sup>, Jan van Zundert<sup>6</sup>, Lennart C. Karszen<sup>2</sup>, Pradeep Suri<sup>5</sup>, Frances M.K. Williams<sup>4</sup>, Yurii S. Aulchenko<sup>3</sup>

<sup>1</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>2</sup>*PolyOmica, 's – Hertogenbosch, the Netherlands*

<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>4</sup>*King's College London, London, UK*

<sup>5</sup>VA Puget Sound Health Care System, Seattle, USA

<sup>6</sup>Maastricht University Medical Centre, Maastricht, The Netherlands

### **Genome – Centered Integrated Instrumental Information System Modeling and Interpretation of Human and Virus Omics**

Anatoliy Shlikht, Natalia Kramorenko

*Far Eastern Federal University, Vladivostok, Russia*

### **The elements of CRISPR – Cas – like system in genome of *Arabidopsis thaliana*: possible origin and some evidence on their functionality**

Ivan Petrushin<sup>1</sup>, Yuri Konstantinov<sup>2</sup>, Igor Gorbenko<sup>2</sup>

<sup>1</sup>*ISU, Irkutsk, Russia*

<sup>2</sup>*SIPPB SB RAS, Irkutsk, Russia*

### **Computational Pipeline for Genomic Epidemiology Surveillance of Pathogenic Bacteria**

Andrey Shelenkov, Yulia Mikhaylova, Yurii Yanushevich, Vasiliy Akimkin

*Central Research Institute of Epidemiology, Moscow, Russia*

### **Genetic mapping of QTLs controlling the ISIAH hypertensive rat behavior in an open field tests**

Olga Redina, Svetlana Smolenskaya, Arcady Markel

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **Transcriptional profiling of ventral tegmental area of male mice with alternative patterns of social behaviors**

Olga Redina, Vladimir Babenko, Vadim Efimov, Dmitry Smagin, Irina Kovalenko, Anna Galyamina, Natalia Kudryavtseva

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **A study of genes controlling carcinogenesis in a regenerative model flatworm *Macrostomum lignano***

Kitill Ustyantsev<sup>1</sup>, Mikhail Biryukov<sup>1</sup>, Eugene Berezikov<sup>1,2</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*European Research Institute for the Biology of Ageing, Groningen, The Netherlands*

### **Novel loci associated with plasma immunoglobulin G N – glycosylation identified by a multivariate analysis**

Alexandra S. Shadrina<sup>1</sup>, Alexander S. Zlobin<sup>1</sup>, Olga O. Zaytseva<sup>2</sup>, Gordan Lauc<sup>2</sup>, Lucija Klaric<sup>3</sup>, Sodbo Z. Sharapov<sup>1</sup>, Yurii S. Aulchenko<sup>1</sup>, Yakov A. Tsepilov<sup>1</sup>

<sup>1</sup>*Institute of Cytology and Genetics, Novosibirsk, Russia*

<sup>2</sup>*Genos Glycoscience Research Laboratory, Zagreb, Croatia*

<sup>3</sup>*University of Edinburgh, Edinburgh, United Kingdom*

### **Peak caller comparison through quality control of ChIP – Seq datasets**

Ruslan N. Sharipov<sup>1,2</sup>, Yury V. Kondrakhin<sup>1,3</sup>, Semyon K. Kolmykov<sup>1,3</sup>, Ivan S. Yevshin<sup>1,3</sup>, Anna S. Ryabova<sup>1,3</sup>, Fedor A. Kolpakov<sup>1,3</sup>

<sup>1</sup>BIOSOFT.RU, LLC; Novosibirsk, Russia

<sup>2</sup>Novosibirsk State University, Novosibirsk, Russia

<sup>3</sup>Institute of Computational Technologies SB RAS, Novosibirsk, Russia

### **The first insights into regulation of cell transdifferentiation during gut regeneration in *Eupentacta fraudatrix***

Alexey Boyko, Igor Dolmatov

NSCMB FEB RAS, Vladivostok, Russia

### **Disruptive natural selection by male reproductive potential prevents underexpression of the genes encoding proteins on the human Y chromosome as a self – domestication syndrome**

Mikhail Ponomarenko, Irina Chadaeva, Dmitry Oshchepkov, Dmitry Rasskazov, Alexander Osadchuk, Ludmila Osadchuk

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **The limits of the additive model for adult height**

Ivan A. Kuznetsov<sup>1,2</sup>, Sergei A. Slavskii<sup>2</sup>, Tatiana I. Shashkova<sup>2</sup>, Georgii A. Bazykin<sup>1</sup>, Tatiana I. Axenovich<sup>3</sup>, Fyodor A. Kondrashov, Yurii S. Aulchenko

<sup>1</sup>Skolkovo Institute of Science and Technology, Moscow, Russia

<sup>2</sup>Novosibirsk State University, Novosibirsk, Russia

<sup>3</sup>Institute of Cytology and Genetics SB RASm Novosibirsk, Russia

<sup>4</sup>Institute of Science and Technology, Vienna, Austria

### **Functional Roles of the E3 Ubiquitin Ligase HYD in *Drosophila* Tissues**

Iuliia Aleksandrovna Galimova<sup>1</sup>, Natalia Vladimirovna Dorogova<sup>2</sup>, Svetlana Aleksandrovna Fedorova<sup>2</sup>, Elena Ustinovna Bolobolova<sup>2</sup>

<sup>1</sup> Institute of Molecular and Cellular Biology, SB RAS, Novosibirsk, Russia

<sup>2</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

### **Competition and collaboration in the miRNA science field**

Artemiy Firsov<sup>1</sup>, Igor Titov<sup>2</sup>

<sup>1</sup>Computer Science and Computer Engineering, Institute of Informatics Systems, Novosibirsk, Russia

<sup>2</sup>The Federal Research Center Institute of Cytology and Genetics, Novosibirsk, Russia

### **High performance pipeline for the calculation of Polygenic Risk Scores**

Arina Nostaeva<sup>1</sup>, Tatiana Shashkova<sup>1</sup>, Sodbo Sharapov<sup>1</sup>, Yakov Tsepilov<sup>1</sup>, Yurii Aulchenko<sup>1,2</sup>, Lennart C. Karssen<sup>2</sup>

<sup>1</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>2</sup>PolyKnomics's – Hertogenbosch, The Netherlands

### **Computer methods for visualization chromosome – specific DNA sequences in FISH images**

Bogomolov A.G.<sup>1,2</sup>, Karamysheva T.V.<sup>1</sup>, Rubtsov N.B.<sup>1,2</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

### **Genome – wide association study of Parkinson’s disease using MAX3 test**

Georgii Ozhegov<sup>1,2</sup>, Dmitry Poveri<sup>3</sup>, Sergey Medvedev<sup>4</sup>, Suren Zakian<sup>4</sup>, Yuri Vyatkin<sup>2,5</sup>, Sergey Postovalov<sup>2,5</sup>

<sup>1</sup>*Kazan Federal University, Kazan, Russia*

<sup>2</sup>*Novel Software Systems, Ltd., Novosibirsk, Russia*

<sup>3</sup>*Novosibirsk State Technical University, Novosibirsk, Russia*

<sup>4</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>5</sup>*Novosibirsk State University, Novosibirsk, Russia*

### **A new method for combining of genetically correlated traits by maximizing of their shared heritability**

Gulnara R. Svishcheva<sup>1</sup>, Evgeny S. Tiys<sup>1</sup>, Sofya G. Feoktistova<sup>1</sup>, Elizaveta E. Elgaeva<sup>1</sup>, Sodbo Sharapov<sup>1</sup>, Yakov A. Tsepilov<sup>2</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

### **Statistical relations between N – glycome of circulating immunoglobuline G and total plasma N – Glycome**

Sofya G. Feoktistova<sup>1</sup>, Tim Spector<sup>2</sup>, Yurii S. Aulchenko<sup>3</sup>, Sodbo Sharapov<sup>3</sup>, Gordan Lauc<sup>4</sup> Yakov A. Tsepilov<sup>5</sup>, Frano Vuckovic<sup>6</sup>

<sup>1</sup>*Institute of Cytology and Genetic, Novosibirsk, Russia*

<sup>2</sup>*Department of Twin Research and Genetic Epidemiology, School of Life Course Sciences King’s College London, London, United Kingdom*

<sup>3</sup>*Institute of Cytology and Genetics, Novosibirsk, Russia*

<sup>4</sup>*Genos Glycoscience Research Laboratory, Zagreb, Croatia*

<sup>5</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>6</sup>*Genos Glycoscience Research Laboratory, Zagreb, Croatia*

### **lncRNAs – their potential in regulation of hypertension and behavior of ISIAH rats**

Ivan Sidorenko, Vladimir Babenko, Arcady Markel, Olga Redina

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **Transcription factor Kaiso regulates cell division in the developing mouse brain**

Nina Illarionova<sup>1</sup>, Maria Borisova<sup>1</sup>, Ekaterina Bazhenova<sup>1</sup>, Daria Fursenko<sup>2</sup>, Daria Zabelina<sup>3</sup>, Nikita Khotskin<sup>1</sup>, Alexander Kulikov<sup>1</sup>

<sup>1</sup>*ICG SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Institute of Gene Biology RAS, Moscow, Russia*

<sup>3</sup>*NSU, Novosibirsk, Russia*

### **Patterns of maternal and paternal inheritance in Russian populations**

Anton Logachev, Daisuke Hirata, Gaik Tamazian

*St. Petersburg State University, St. Petersburg, Russia*

### **Molecular basis of phosphoryl guanidine oligonucleotides elongation by Taq DNA polymerase**

Alexander Lomzov, Dmitrii Pyshnyi

*ICBFM SB RAS, Novosibirsk, Russia*

### **Software pipeline for the analysis of the functional role of nucleotide substitutions in regulatory regions of genes and its testing on polymorphisms associated with obesity**

Ekaterina Alekseevna Matrosova<sup>1</sup>, Vadim Mikhailovich Efimov<sup>1,2</sup>, Elena Vasilevna Ignatieva<sup>2</sup>

<sup>1</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **WebMCOT web – service for prediction of co – occurred DNA motifs in ChIP – seq data**

Aleksey Mukhin, Victor Levitsky, Dmitriy Y. Oschepkov, Sergey A. Lashin

*Institute Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **Identification and description of the genes with a high mutation frequency in vagal paragangliomas**

Vladislav Pavlov<sup>1</sup>, Anastasiya Snezhkina<sup>1</sup>, George Krasnov<sup>1</sup>, Dmitry Kalinin<sup>2</sup>, Alexander Golovyuk<sup>2</sup>, Anna Kudryavtseva<sup>1</sup>

<sup>1</sup>*EIMB RAS, Moscow, Russia*

<sup>2</sup>*A.V. Vishnevsky National Medical Research Center of Surgery, Moscow, Russia*

### **Differentially expressed microRNAs in carotid paraganglioma**

Anastasiya Snezhkina, Elena Pudova, Vladislav Pavlov, Maria Fedorova, George Krasnov, Anna Kudryavtseva

*EIMB RAS, Moscow, Russia*

### **Results of the whole – genomic sequencing and annotation of the Listeria phenotype**

Marina Terekhova<sup>1</sup>, Elizaveta Rogacheva<sup>2</sup>, Lyudmila Kraeva<sup>2</sup>, Irina Derevyanchenko<sup>3</sup>

<sup>1</sup>*St. Petersburg State University of Information Technologies, Mechanics and Optics, Saint – Petersburg, Russia*

<sup>2</sup>*Saint – Petersburg Pasteur Institute, Saint – Petersburg, Russia*

<sup>3</sup>*Branch of the Federal State Health Institution “Center for Hygiene and Epidemiology in the City of St. Petersburg”, Saint – Petersburg, Russia*

### **LTR – retrotransposon transcripts are ubiquitously expressed, polyadenylated and underwent splicing in sunflower (*Helianthus annuus* L.)**

Pavel Merkulov, Murad Omarov, Ilya Kirov

*All – Russian Research Institute of Agricultural Biotechnology RAS, Moscow, Russia*

### **Constructing a pipeline for genome variant / gene functioning hybrid prioritization: a case study of type II diabetes**

Irina Kolesnikova<sup>1</sup>, Valery Polunovsky<sup>1</sup>, Konstantin Gunbin<sup>2,3</sup>

<sup>1</sup>LLC NCGI, Novosibirsk, Russia

<sup>2</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>3</sup>NSU, Novosibirsk, Russia

### **Differentially expressed genes associated with TMPRSS2 – ERG molecular subtype of prostate cancer**

Anastasiya Andreevna Kobelyatskaya<sup>1</sup>, Elena Anatolevna Pudova<sup>1</sup>, George Sergeevich Krasnov<sup>1</sup>, Anna Victorovna Kudryavtseva<sup>1</sup>, Kirill Mikhailovich Nyushko<sup>2</sup>, Boris Yakovlevich Alekseev<sup>2</sup>

<sup>1</sup>EIMB RAS, Moscow, Russia

<sup>2</sup>FSBI NMRRC, Moscow, Russia

### **RTrans: a pipeline for multi – way analysis of differential gene expression profiles**

George Sergeevich Krasnov, Anastasiya Andreevna Kobelyatskaya, Anastasiya Vladimirovna Snezhkina, Vladislav Sergeevich Pavlov, Elena Anatolevna Pudova, Anna Victorovna Kudryavtseva

EIMB RAS, Moscow, Russia

### **Intermediate and high – risk prostate cancer methylation analysis**

Anastasiya Andreevna Kobelyatskaya<sup>1</sup>, Kirill Mikhailovich Nyushko<sup>2</sup>, Elena Anatolevna Pudova<sup>1</sup>, Boris Yakovlevich Alekseev<sup>2</sup>, George Sergeevich Krasnov<sup>1</sup>, Anna Victorovna Kudryavtseva<sup>1</sup>

<sup>1</sup>EIMB RAS, Moscow, Russia

<sup>2</sup>FSBI NMRRC, Moscow, Russia

### **Allelic drop – out is a common phenomenon reducing the diagnostic yield of PCR – based target sequencing**

Anna Shestak<sup>1</sup>, Anna Bukaeva<sup>1</sup>, Siamak Saber<sup>2</sup>, Elena Zaklyazminskaya<sup>1</sup>

<sup>1</sup>Petrovsky National Research Center of Surgery, Moscow, Russia

<sup>2</sup>Cardiac Electrophysiology Research Center, Rajaie Cardiovascular Medical and Research Center, Iran University of Medical Sciences, Tehran, Iran

### **AD ASTRA: the database of Allelic Dosage – corrected Allele – Specific TRAnscription factor binding suggests causal regulatory sequence variants of pathologies**

Sergey Abramov<sup>1</sup>, Alexandr Boytsov<sup>1</sup>, Bykova Dariia<sup>2</sup>, Eugene Baulin<sup>3</sup>, Ivan Yevshin<sup>4</sup>, Fedor Kolpakov<sup>6</sup>, Vsevolod J Makeev<sup>1</sup>, Ivan V Kulakovskiy<sup>5</sup>

<sup>1</sup>Vavilov Institute of General Genetics Russian Academy of Sciences, Moscow, Russia

<sup>2</sup>Lomonosov Moscow State University, Moscow, Russia

<sup>3</sup>Institute of Mathematical Problems of Biology RAS – the Branch of Keldysh Institute of Applied Mathematics of Russian Academy of Sciences, Pushchino, Russia

<sup>4</sup>BIOSOFT.RU LLC, Novosibirsk, Russia

<sup>5</sup>Engelhardt Institute of Molecular Biology, Moscow, Russia

<sup>6</sup>Institute of Computational Technologies SB RAS, Novosibirsk, Russia

### **Analysis of short – and long – range interactions within potential binding sites notably extends the fraction of verified peaks in ChIP – seq data**

Anton Tsukanov, Victor Levitsky, Tatyana Merkulova

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **X-chromosome Inactivation in American Mink iPSCs**

Inna Pristyazhnyuk Aleksei Menzorov

*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

### **Cryptic Plasmids of Alfalfa Root Nodule Bacteria – Structural and Functional Diversity**

Alla Saksaganskaia, Viktoria Muntyan, Alexey Afonin, Marina Roumiantseva

*ARRIAM, Saint – Petersburg, Pushkin, Russia*

### **Whole genome sequencing and assembly of *Saccharomyces cerevisiae* genomes using Oxford Nanopore data**

Andrew G. Matveenko<sup>1</sup>, Anton B. Matiiv<sup>1</sup>, Yury A. Barbitoff<sup>1,4</sup>, Evgenia M. Maksiutenko<sup>1,2</sup>, Svetlana E. Moskalenko<sup>1,2</sup>, Alexandra V. Beliavskaia<sup>3</sup>, Alexander V. Predeus<sup>3,4</sup>, Galina A. Zhouravleva<sup>1</sup>

<sup>1</sup>*St. Petersburg State University, St. Petersburg, Russia*

<sup>2</sup>*Vavilov Institute of General Genetics, St. Petersburg Branch, St. Petersburg, Russia*

<sup>3</sup>*University of Liverpool, Liverpool, UK*

<sup>4</sup>*Bioinformatics Institute, St. Petersburg, Russia*

### **The Rich Inner World of Colorado Potato Beetles – a Metagenomic Survey of Viral Diversity in Public Data**

Maria Starchevskaya<sup>1</sup>, Yuri Vyatkin<sup>2,3</sup>, Denis Antonets<sup>1,3</sup>

<sup>1</sup>*SRC VB “Vector” Rospotrebnadzor, Koltsovo, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*Novel Software Systems LLC, Novosibirsk, Russia*

### **Potential of Whole Genome Sequencing in the Assessment of Sensitivity of Clinical Isolate *M. tuberculosis* to Antibiotics**

Olga Berdyugina

*IIP RAS, Ekaterinburg, Russia*

### **Genome distance between regulatory elements of growth – related genes may determine morpho – physiological traits in mammals**

Dmitriy Romanov, Tatiana Shkurat

*Southern federal university, Rostov – on – Don, Russia*

### **Promoter expression landscape in skeletal muscle in hindlimb suspension and recovery model in rat**

Guzel Gazizova<sup>1</sup>, Ruslan Deviatiiarov<sup>1</sup>, Islam Nigmatzhanov<sup>1</sup>, Ilia Akberdin<sup>2,3</sup>, Sergei Pintus<sup>2,4</sup>, Oksana Tyapkina<sup>5</sup>, Fedor Kolpakov<sup>4</sup>, Leniz Nurullin<sup>5</sup>, Oleg Gusev<sup>5,6</sup>

<sup>1</sup>*KFU, Kazan, Russia*

<sup>2</sup>*BIOSOFT.RU, LLC;*

<sup>3</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>4</sup>*Institute of Computational Technologies, Novosibirsk, Russia*

<sup>5</sup>*FRC KSC RAS, Kazan, Russia*

<sup>6</sup>*RIKEN, Yokohama, Japan*

### **Differentially methylation of ANKRD53 and GATA3 genes in human miscarriages with trisomy 16**

E. N. Tolmacheva | S.A.Vasilyev | O.Yu. Vasilyeva | T.V. Nikitina | E.A.Sazhenova | A.V.Markov | E.S. Serdyukova | D.I. Zhigalina | I.N.Lebedev

### **Advanced data curation in GTRD database: hierarchical dictionaries of cell types and experimental factors**

Mikhail A. Kulyashov<sup>1,2,3,4</sup>, Semyon K. Kolmykov<sup>1,3,4</sup>, Ivan S. Yevshin<sup>1,4</sup>, Fedor A. Kolpakov<sup>1,4</sup>

<sup>1</sup>*BIOSOFT.RU, LLC, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*FRC Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>4</sup>*Institute of Computational Technologies SB RAS, Novosibirsk, Russia*

### **Functional annotation of the transcription factors from *Methylotuvimicrobium alcaliphilum* 20ZR**

Semyon K. Kolmykov<sup>1,2,3</sup>, Nikita V. Ivanisenko<sup>3</sup>, Ivan S. Evshin<sup>1,2</sup>, Mikhail Kulyashov<sup>1,2,4</sup>, Tamara M. Khlebodarova<sup>3</sup>, Ilya R. Akberdin<sup>1,3,4</sup>

<sup>1</sup>*BIOSOFT.RU, LLC, Novosibirsk, Russia*

<sup>2</sup>*Institute of Computational Technologies, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*FRC Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>4</sup>*Novosibirsk State University, Novosibirsk, Russia*

### **Transcriptome (RNA – seq) analysis of human salivary gland cells with exogenous expression of human pancreas beta cells transcription factors PDX1, MAFA, NGN3**

Olga Brovkina<sup>1</sup>, Alexander Artyuhov<sup>2</sup>, Yulia Kolesova<sup>3</sup>, Erdem Dashinimaev<sup>2,4</sup>, Mikhail Borisov<sup>4</sup>, Ekaterina Vorotelyak<sup>4</sup>, Andrey Vasiliev<sup>4</sup>

<sup>1</sup>*Federal Research and Clinical Center, FMBA of Russia, Moscow, Russia*

<sup>2</sup>*Pirogov Russian National Research Medical University, Moscow, Russia*

<sup>3</sup>*Sechenov First Moscow State Medical University, Institute of Molecular Medicine, Moscow, Russia*

<sup>4</sup>*Koltzov Institute of Developmental Biology, Russian Academy of Sciences, Moscow, Russia*

### **Metavirome analysis of Baikal sponges**

Tatyana Vladimirovna Butina, Yuriy Sergeevich Bukin, Igor Veniaminovich Khanaev

*LIN SB RAS, Irkutsk, Russia*

### **Detection of alphacoronavirus in bat fecal samples from Volgograd region**

Elena Korneenko<sup>1</sup>, A.E. Samoilov<sup>1</sup>, I.V.Artyushin<sup>2</sup>, A.V. Dudorova<sup>2</sup>, E.V. Pimkina<sup>1</sup>, V.G. Dedkov<sup>3</sup>, M.V. Safonova<sup>4</sup>, A.D. Matsvay<sup>5</sup>, A.S. Speranskaya<sup>1,2</sup>

<sup>1</sup>*Central Research Institute of Epidemiology, Moscow, Russia*

<sup>2</sup>*Lomonosov Moscow State University, Moscow, Russia*

<sup>3</sup>*Pasteur Institute, Saint – Petersburg, Saint – Petersburg, Russia*

<sup>4</sup>*Plague Control Center, Federal Service on Consumers' Rights Protection and Human Well – Being Surveillance, Moscow, Russia*

<sup>5</sup>*FSBI “Center of Strategic Planning” of the Ministry of Health, Moscow, Russia*

### **Genome – wide Association Study Reveals Novel Genetic Variants Associated with HIV – 1C Infection in Botswana Population**

Andrey Shevchenko, Sergey V. Malov, Alexey Antonik

*Theodosius Dobzhansky Center for Genome Bioinformatics, St – Petersburg, Russia*

*St. – Petersburg State University, St – Petersburg, Russia*

### **Automatic Annotation of Operons Responsible for O – antigen Synthesis**

Danil Zilov, Polina Chesnokova, Alexey Komissarov

*ITMO University, St. Petersburg, Russia*

### **Short sequence repeats (SSR) under selection pressure: Cyprinidae fish case study**

Mikhail Orlov<sup>1</sup>, Andrey Tykhonov<sup>2</sup>

<sup>1</sup>*ICB RAS, Pushchino, Russia*

<sup>2</sup>*“Aqua Logo” company group, Moscow, Russia*

### **Evaluation of biological activity of the conjugates of granulocyte – macrophage colony stimulating factor with alendronic acid**

Ekaterina A. Volosnikova, Tat'yana I. Esina, Alena V. Bateneva, Galina G. Shimina, Elena D. Danilenko

*IMBT FBRI SRC VB «Vector», Rospotrebnadzor, Berdsk, RUSSIA*

### **Functional study of potential regulatory SNPs (rs590352, rs11542583, rs3829202, rs78317230, rs2072580, rs4796672)**

Arina Degtyareva<sup>1</sup>, Elena Leberfarb<sup>1</sup>, Ilya Brusentsov<sup>1</sup>, Tatiana Kuzina<sup>2</sup>, Tatiana Merkulova<sup>1</sup>

<sup>1</sup>*ICG SB RAS;* <sup>2</sup>*NSU, Novosibirsk, RUSSIA*

### **PCR dependent biases could significantly affect quantitative estimation of plant mix composition**

Valeriia Kaptelova<sup>1</sup>, Maria Logacheva<sup>4,5</sup>, Anna Speranskaya<sup>1</sup>, Denis Omelchenko<sup>2</sup>, Anna Fedotova<sup>4,5</sup>, Anastasia Krinitsina<sup>5,6</sup>, Andrey Ayginin<sup>3</sup>, Kamil Khafizov<sup>3</sup>, Elena Korneenko<sup>1</sup>, Andrei Samoilov<sup>1</sup>

<sup>1</sup>Central Research Institute of Epidemiology, Moscow, Russia

<sup>2</sup>Institute of Information Transmission Problems, Moscow, Russia

<sup>3</sup>Center for Strategic Planning and Management of Biomedical Health Risks, Moscow, Russia

<sup>4</sup>Skolkovo Institute of Science and Technology, Skolkovo, Russia

<sup>5</sup>Lomonosov Moscow State University, Moscow, Russia

<sup>6</sup>I.M. Sechenov First Moscow State Medical University, Pharmaceutical Natural Science Department, Moscow, Russia

### **Bioinformatic Screening for Subtilisin – like Peptidases in Dikaryotic Fungi**

Nikita Alkin<sup>1</sup>, Yakov Dunaevsky<sup>2</sup>, Mikhail Belozersky<sup>2</sup>, Galina Beliakova<sup>1</sup>, Valeriia Tereshchenkova<sup>1</sup>, Elena Elpidina<sup>2</sup>

<sup>1</sup>MSU, Moscow, Russia

<sup>2</sup>MSU Belozersky IPCB, Moscow, Russia

### **EPhIMM: computational workflow for fast phylogenetic inference based on multiple alignment of prokaryotic single – copy marker genes**

Aleksei Korzhenkov

NRC Kurchatov Institute, Moscow, Russia

### **Functioning of unique nitrile – detoxifying system in soil xenobiotic degrader *Rhodococcus rhodochrous*: a whole – genome transcriptomic approach**

Konstantin V Lavrov<sup>1</sup>, Andrey D Novikov<sup>1</sup>, Tatyana I Kalinina<sup>1</sup>, Artem S Kasianov<sup>2</sup>, Alexander S Yanenko<sup>1</sup>

<sup>1</sup>NRC “Kurchatov institute –GosNIIgenetika, Moscow, Russia

<sup>2</sup>Vavilov Institute of General Genetics, Moscow, Russia

### **Comparative analysis of repeatome composition of four allopolyploid Poaceae species**

Elizaveta Kolganova, Michail Divashuk, Ilya Kirov

All – Russia Research Institute of Agriculture Biotechnology, Moscow, Russia

### **Comparison of Brain Transcriptome Profiles of Short – lived and Long – lived Species of *Nothobranchius***

Zulfiia Guvatova<sup>1</sup>, George Krasnov<sup>1</sup>, Sergey Simanovsky<sup>2</sup>, Alexander Frolov<sup>2</sup>, Nataliya Gladyshev<sup>3</sup>, Anna Kudryavtseva<sup>2</sup>

<sup>1</sup>Engelhardt Institute of Molecular Biology, Russian Academy of Sciences, Moscow, Russia

<sup>2</sup>A.N. Severtsov Institute of Ecology and Evolution, Moscow, Russia

<sup>3</sup>MSAVM&B – MVA named after K.I. Skryabin, Moscow, Russia

### **Genome Assembly and Annotation of *Nothobranchius rachovii* killifish**

Zulfiia Guvatova<sup>1</sup>, George Krasnov<sup>1</sup>, Anastasiya Snezhkina<sup>1</sup>, Artemy Tokarev<sup>2</sup>, Maria Fedorova<sup>1</sup>, Anna Kudryavtseva<sup>1</sup>

<sup>1</sup>Engelhardt Institute of Molecular Biology, Russian Academy of Sciences, Moscow, Russia

<sup>2</sup>MSAVM&B – MVA named after K.I. Skryabin, Moscow, Russia

### **Comparative genomics and quantitative proteomics reveal differentially produced proteins underlying virulence and host specificity in *Bacillus thuringiensis***

Yury Malovichko<sup>1</sup>, Maria Belousova<sup>1</sup>, Elena Lukasheva<sup>2</sup>, Daria Gorbach<sup>2</sup>, Ekaterina Romanovskaya<sup>2</sup>, Christian Ihling<sup>3</sup>, Andrej Frolov<sup>2,3</sup>, Anton Nizhnikov<sup>1,2</sup>, Kirill Antonets<sup>1,2</sup>

<sup>1</sup>All – Russian Research Institute of Agricultural Microbiology, Saint Petersburg, Russia

<sup>2</sup>St. Petersburg State University, Saint Peterburg, Russia

<sup>3</sup>Institute of Pharmacy, Martin – Luther Universität Halle – Wittenberg, Halle, Germany

### **Expression of DNA repair genes in anhydrobiotic insect *Polypedilum vanderplanki***

Alexander Nesselov, Sabina Kondratyeva, Taisiya Voronina

IFMB KFU, Kazan, Russia

### **Predicting elongation efficiency of gene translation for annotation of bacterial genomes: a case study for biosynthetic gene clusters of nonribosomal peptides**

A.I. Klimenko<sup>1</sup>, Yu.G. Matushkin<sup>1</sup>, D.A. Afonnikov<sup>1,2</sup>

<sup>1</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>2</sup>Novosibirsk State University, Novosibirsk, Russia

### **MicroRNA content of horse and human milk exosomes**

Sergey Sedykh, Kuleshova Anna, Georgy Nevinsky

ICBFM SB RAS, Novosibirsk, Russia

### **Mutational profile of Diffuse Large B – cell Lymphoma with central nervous system relapse: analysis of CBioPortal for Cancer Genomics database**

Elena Voropaeva<sup>1</sup>, Olga Beresina<sup>3</sup>, Viktoria Karpova<sup>4</sup>, Yuriy Orlov<sup>2</sup>, Maria Churkina<sup>3</sup>, Tatyana Pospelova<sup>3</sup>, Vladimir Maximov<sup>1</sup>, Anastasia Ivanova<sup>1</sup>, Elizaveta Melnikova<sup>1</sup>, Anna Gurageva<sup>1</sup>

<sup>1</sup>IITPM – Branch of Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>2</sup>I.M. Sechenov First Moscow State Medical University, Moscow, Russia

<sup>3</sup>NSMU, Novosibirsk, Russia

<sup>4</sup>State Regional Clinical Hospital, Novosibirsk, Russia

### **Justification of measures for optimization and prevention with dysplasia of stratified squamous epithelium of the cervix in women of reproductive age**

Ra'nokhon Solieva Bakhodir qizi<sup>1</sup>, Dilfuza Alieva Abdullaevna<sup>2</sup>

<sup>1</sup>ASMI, Andijan, Uzbekistan

<sup>2</sup>Republican specialized scientific and practical medical center of obstetrics and gynecology. Tashkent, Uzbekistan

### **Hemolymph metagenome of endemic amphipod *Eulimnogammarus verrucosus* from Lake Baikal**

Ekaterina Shchapova<sup>1</sup>, Anton Gurkov<sup>1</sup>, Natalia Belkova<sup>2</sup>, Renat Adelshin<sup>3,1</sup>, Maxim Timofeyev<sup>1</sup>

<sup>1</sup>*Irkutsk State University, Irkutsk, Russia*

<sup>2</sup>*Scientific Centre for Family Health and Human Reproduction Problems, Irkutsk, Russia*

<sup>3</sup>*Irkutsk Anti – Plague Research Institute of Siberia and Far East, Irkutsk, Russia*

### **Genes expression related to the effects of hypoxia in the marine mussel, *Mytilus galloprovincialis***

Ekaterina Vodiasova<sup>1</sup>, Aleksandra Andreyeva<sup>1</sup>, Anastasiya Lantushenko<sup>2</sup>, Yakov Meger<sup>2</sup>, Irina Deghtyar<sup>2</sup>, Dmitry Afonnikov<sup>3,2</sup>

<sup>1</sup>*IBSS RAS, Sevastopol, Russia*

<sup>2</sup>*SSU, Sevastopol, Russia*

<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

## SYMPOSIUM

# Systems computational biology: analysis, mathematical modeling and information technologies

**Oral reports**

7 July, Tuesday

**Small Conference Hall**

**Morning session. Systems computational biology: analysis, mathematical modeling and information technologies**

**Chair: Mikhail Marchenko, Institute of Computational Mathematics and Mathematical Geophysics, SB RAS, Novosibirsk, Russia  
Denis Ponomarev, A.P. Ershov Institute of Informatics Systems SB RAS, Novosibirsk, Russia**

09:00 – 09:30

**PLENARY REPORT**

**Understanding heterozygous carrying of deleterious variants as a key for personalized longevity**

*Ancha Baranova,*

*School of Systems Biology, George Mason University, Fairfax, VA*

9:30 – 10:00

**Keynote report**

**ANDSystem: text mining – based associative gene networks discovery system and its application to biomedical tasks**

*Vladimir Ivanisenko<sup>1,2</sup>, Olga Saik<sup>1</sup>, Timofey Ivanisenko<sup>1,2</sup>, Nikita Ivanisenko<sup>1</sup>, Evgeny Tiys<sup>1</sup>, Pavel Demenkov<sup>1</sup>, Nikolay Kolchanov<sup>1</sup>*

*<sup>1</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

*<sup>2</sup>Novosibirsk State University, Novosibirsk, Russia*

10:00 – 10:20

**ANDDigest: A Text – Mining Based Computer System For Generating Digests in the Field of Biology**

*Timofey Ivanisenko<sup>1,2</sup>, Pavel Demenkov<sup>1</sup>, Vladimir Ivanisenko<sup>1,2</sup>, Nikolay Kolchanov<sup>1</sup>*

*<sup>1</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

*<sup>2</sup>Novosibirsk State University, Novosibirsk, Russia*

10:20 – 10:40

**A feedback loop enrichment analysis in gene network of Bronchial asthma and pulmonary tuberculosis interaction**

*Evgeny S. Tiys, Pavel S. Demenkov, Vladimir A. Ivanisenko, Nikolay A. Kolchanov*

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

10:40 – 11:00

**Integrated informational – computer system for modeling and analysis of DNA functional sites activity**

*Mikhail Ponomarenko*

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

11:00 – 11:20

**Coffee – break**

11:20 – 11:40

**The novel primary targets of CDDO – Im, defining its cytoprotective activity:**

***in silico* identification**

*Andrey Markov*

*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

11:40 – 12:00

**Analysis of noise in gene ensembles based on transcriptional responses of the human body to coronavirus infection: a search for predictors of infection severity, level of immune response, and new pharmacological targets**

*Yu. M. Moschkin*

*Erasmus MC, Rotterdam, Netherlands*

12:00 – 12:20

**Mathematical Modeling of Allergenic Pollen Propagation in Atmospheric Layer**

*Olga Sergeevna Medveditsyna<sup>1</sup>, Sergey Leonidovich Rychkov<sup>2</sup>, Anatoly Victorovich Shatrov<sup>2</sup>*

*<sup>1</sup>Kirov State Medical University, Kirov City, Russia*

*<sup>2</sup>Vyatka State University, Kirov City, Russia*

12:20 – 12:40

**On region based inference in genome wide association study**

Sergey V. Malov, Alexey Antonik, Andrey Shevchenko

*Theodosius Dobzhansky Center for Genome Bioinformatics, St – Petersburg, Russia*

*St. – Petersburg State University, St – Petersburg, Russia*

Lunch

**Evening session 1. Mathematical issues of systems biology**

**Chairs: Vladimir Golubyatnikov, *Sobolev Institute of Mathematics, SB RAS, Novosibirsk, Russia*;**

**Andrey Palyanov, *A.P. Ershov Institute of Informatics Systems, SB RAS, Novosibirsk, Russia***

15.00 – 15.30

**Keynote report**

**Mathematics of Covid – 19**

S.I. Kabanikhin, O.I. Krivorotko, A.Yu. Prikhodko, N.M. Prokhoshin, M.A. Shishlenin, N.Yu. Zyatkov

*The Institute of Computational Mathematics and Mathematical Geophysics, SB RAS, Novosibirsk, Russia*

15.30 – 16.00

**Runtime analysis of non – elitist evolutionary algorithms with fitness – proportionate selection on Royal Road functions**

Anton Eremeev

*The Institute of Scientific Information for Social Sciences, RAS, Moscow, Russia*

*Omsk Branch of Sobolev Institute of Mathematics, Omsk, Russia*

16.00 – 16.30

**Phase Portraits of Gene Networks Models**

Natalia Ayupova<sup>1</sup>, Vladimir Golubyatnikov<sup>1</sup>, Vyacheslav Gradov<sup>2</sup>, Liliya Minushkina<sup>2</sup>

<sup>1</sup>*Sobolev Institute of Mathematics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University Novosibirsk, Russia*

16.30 – 16.50

**Adjoint Ensemble Methods for Inverse Modeling of Biological Processes**

Alexey Penenko<sup>1</sup>, Ulyana Zubairova<sup>2</sup>, Alexey Doroshkov<sup>2</sup>, Alexander Bobrovskikh<sup>2</sup>

<sup>1</sup>*Institute of Computational Mathematics and Mathematical Geophysics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

16.50 – 17.10

Coffee – break

17.10 – 17.30

**Stability of equilibrium points in a predator – prey model with delayed argument**

Maria Skvortsova, Timur Yskak

*Sobolev Institute of Mathematics, Novosibirsk, Russia*

**17.30 – 17.50 Digital Platform “Bioinformatics”: System – Forming Solutions**

Yurii Zybarev, [Sergey Kratov](#)

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**17.50 – 18.10 Anchored Bootstrap**

[Vadim Efimov](#)<sup>1,2</sup>, Kirill Efimov<sup>3</sup>, Vera Kovaleva<sup>4</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*IHNA&NPh RAS, Moscow, Russia*

<sup>4</sup>*ISEA SB RAS, Novosibirsk, Russia*

**18.10 – 18.30 Autoencoder – based Lowdeg – Rank Spectral Ensemble Clustering of Biological Data**

Vladimir Berikov

*Sobolev Institute of Mathematics, SB RAS, Novosibirsk, Russia*

**18.30 – 18.50 Multi – class brain tumor segmentation via multi – sequences MRI mixture data preprocessing.**

Andrey Letyagin<sup>1</sup>, Sergey Golushko<sup>2</sup>, Mikhail Amelin<sup>3</sup>, Bair Tuchinov<sup>2</sup>, Evgeniya Amelina<sup>2</sup>, Nikolay Tolstokulakov<sup>2</sup>, Evgeniy Pavlovskiy<sup>2</sup>, Vladimir Groza<sup>4</sup>

<sup>1</sup>*Research Institute of Clinical and Experimental Lymphology, Branch of IC&G SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*FSBI “Federal Neurosurgical Center”, Novosibirsk, Russia*

<sup>4</sup>*Median Technologies, Valbonne, France*

## 8 July, Wednesday

### Computer Class

### Evening Session 2. Mathematical issues of systems biology

**Chair: Matteo Barberis, University of Surrey Guildford, Surrey, United Kingdom;  
S.A. Lashin, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia**

**14.30 – 15.00**

**PLENARY REPORT**

**Understanding the evolution of complex regions in bird genomes**

Alexander Sang-Jae Suh,

*Department of Evolutionary Biology, Uppsala University, Uppsala, Sweden*

15.00 – 15.30

**Keynote report**

**A computational approach to investigation of *C. elegans* backwards crawling mechanism via simulation of involved nervous and muscular cells activity driving body movement**

Andrey Yu. Palyanov<sup>1,2</sup>, Natalia V. Palyanova<sup>3</sup>

<sup>1</sup>*A.P. Ershov Institute of Informatics Systems, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*Institute of Molecular Biology and Biophysics, Novosibirsk, Russia*

15.30 – 16.00

**Keynote report**

**Software frameworks for modeling complex hierarchical biological systems**

S.A. Lashin

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

16.00 – 16.10

**MGSGenerator 1.5: software tool for reconstructing mathematical models of metabolic networks**

F.V. Kazantsev<sup>1</sup>, S.A. Lashin<sup>1,2</sup>

<sup>1</sup>*Kurchatov genomics center Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

16.10 – 16.30

**Motility and fitness of microorganisms in dynamic aquatic ecosystems: a simulation study**

A.I. Klimenko, Yu.G. Matushkin, S.A. Lashin

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

16.30 – 16.50

**BioUML – universal platform for analyses of biomedical data**

Fedor A. Kolpakov<sup>1,2</sup>, Anna S. Ryabova<sup>1,2</sup>, Elena O. Kutumova<sup>1,2</sup>, Ivan S. Evshin<sup>1,2</sup>, Yury V. Kondrakhin<sup>1,2</sup>, Nikita V. Mandrik<sup>1,2</sup>, Ilya N. Kiselev<sup>1,2</sup>, Sergey S. Pintus<sup>1,2</sup>, Alexander E. Kel<sup>2,3</sup>

<sup>1</sup>*Institute of Computational Technologies SB RAS, Novosibirsk, Russia*

<sup>2</sup>*BIOSOFT.RU, LLC, Novosibirsk, Russia*

<sup>3</sup>*geneXplain GmbH, Wolfebuttel, Germany*

16.50 – 17.10

**Coffee – break**

17.10 – 17.30

**Systems biology analysis of metabolism, signaling and gene expression regulation in human skeletal muscle**

Ilya R. Akberdin<sup>1,2,3</sup>, Alexander Yu. Vertyshev<sup>4</sup>, Ilya N. Kiselev<sup>1,5</sup>, Pavel A. Makhnovskii<sup>6</sup>, Fedor A. Kolpakov<sup>1,5</sup>, Sergey S. Pintus<sup>1,5</sup>, Daniil V. Popov<sup>6</sup>

<sup>1</sup>*BIOSOFT.RU, LLC, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>4</sup>CJSC "Sites – Tsentr" Moscow, Russia

<sup>5</sup>Institute of Computational Technologies SB RAS, Novosibirsk, Russia

<sup>6</sup>Institute of Biomedical Problems of the RAS, Moscow, Russia

17.30 – 17.50

**Genome – scale metabolic modeling of 2,3 – butanediol production by *Geobacillus icigianus***

Mikhail Kulyashov<sup>1,2,3</sup>, Ilya R. Akberdin<sup>1,3,4</sup>

<sup>1</sup>BIOSOFT.RU, LLC, Novosibirsk, Russia

<sup>2</sup>Institute of Computational Technologies SB RAS, Novosibirsk, Russia

<sup>3</sup>Novosibirsk State University, Novosibirsk, Russia

<sup>4</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

17.50 – 18.10

**Modeling the mutual relationship between the circadian clock and inflammation response**

Nikolay Podkolodnyy, Natalya Tverdokheb, Olga Podkolodnaya

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

18.10 – 18.30

***In silico* model of glioma MTS growth. Effects of compression and mechanical ECM remodeling**

Vladimir Kalinin

*R&D Sector of TMA, Dundalk, Ireland*

18.30 – 18.50

**SINE and LINE – 1 competition for energy resources determines cell fate**

Maria Duk<sup>1</sup>, Alexandra Chertkova<sup>2,3</sup>, Vitaly Gursky<sup>1,2</sup>, Maria Samsonova<sup>2</sup>, Alexander Kanapin<sup>4</sup>, Anastasia Samsonova<sup>4</sup>

<sup>1</sup>*The Ioffe Institute, St. Petersburg, Russia; St. Petersburg University, St.Petersburg, Russia*

<sup>2</sup>*Peter the Great St.Petersburg Polytechnic University, St.Petersburg, Russia*

<sup>3</sup>*BioCAD, St.Petersburg, Russia*

<sup>4</sup>*St. Petersburg State University, St.Petersburg, Russia*

## Poster session

### Session 1. Systems computational biology

#### A Model of one Central Regulatory Circuit

Tatyana Bukharina<sup>1</sup>, Andrey Akinshin<sup>2</sup>, Vladimir Golubyatnikov<sup>2</sup>, Dagmara Furman<sup>1,3</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Sobolev Institute of Mathematics, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Novosibirsk State University, Novosibirsk, Russia*

### **Development of a method for recognizing biomedical entities in the texts of scientific articles**

Stepan Derevyanchenko<sup>1</sup>, Pavel Demenkov<sup>2</sup>

<sup>1</sup>Novosibirsk State University Novosibirsk, Russia

<sup>2</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, **Russia**

### **Mathematical model of punctuated equilibrium evolution**

Vitaly A. Likhoshvai, Tamara M. Khlebodarova

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **Gene Network of Type 2 Diabetes: Reconstruction and Analysis**

Vladimir Zamyatin<sup>1,2</sup>, Dmitry Afonnikov<sup>1,2</sup>, Zakhar Mustafin<sup>1,2</sup>, Vadim Klimontov<sup>1,2</sup>,

Yury Matushkin<sup>1,2</sup>, Sergey Lashin<sup>1,2</sup>

<sup>1</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>2</sup>Novosibirsk State University, Novosibirsk, Russia

### **Integration of transcriptomics data into a genome – scale metabolic model of the methanotrophic bacterium *Methylothermobacter alkaliphilum* 20Z<sup>R</sup>**

Mikhail Kulyashov<sup>1,2,3,4</sup>, Semyon K. Kolmykov<sup>1,2,4</sup>, Ivan S. Evshin<sup>1,3</sup>, Tamara M. Khlebodarova<sup>2</sup>, Nikita V. Ivanisenko<sup>2</sup>, Ilya R. Akberdin<sup>1,2,4</sup>

<sup>1</sup>BIOSOFT.RU, LLC, Novosibirsk, Russia

<sup>2</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>3</sup>Institute of Computational Technologies, SB RAS, Novosibirsk, Russia

<sup>4</sup>Novosibirsk State University, Novosibirsk, Russia

### **Transfer of Genetic Algorithms to Directed Evolution of Macromolecules: Tests *in silico***

Ekaterina Myasnikova<sup>1</sup>, Alexander Spirov<sup>2</sup>

<sup>1</sup>Peter the Great St. Petersburg Polytechnical University Saint – Petersburg, Russia

<sup>2</sup>I.M. Sechenov Institute of Evolutionary Physiology and Biochemistry Russian Academy of Sciences Saint – Petersburg, Russia

### **ECM stiffness effects and subumor formation in glioma growth. *In silico* model**

Vladimir Kalinin

*R&D Sector of TMA, Dundalk, Ireland*

### **Development and analysis of AIDS epidemic agent – based computer model applying an algorithm for explicit calculation of HIV replicability**

Anna Smirnova<sup>1,2</sup>, Mikhail Ponomarenko<sup>1</sup>, Sergey Lashin<sup>1,2</sup>

<sup>1</sup>ICG SB RAS, Novosibirsk, Russia

<sup>2</sup>NSU, Novosibirsk, Russia

## Session 2. Mathematical issues of systems biology

### Named entity recognition in medical texts in Russian using deep learning models

*I.V. Moskalev, L.A. Khvorova*  
*ASU, Barnaul, Russia*

### The algorithm for finding potentially oscillating behavior in enzymatic systems

*Tatiana N. Lakhova*<sup>1</sup>, *Fedor V. Kazantsev*<sup>1</sup>, *Yuriy G. Matushkin*<sup>2</sup>, *Sergey A. Lashin*<sup>1,3</sup>

<sup>1</sup>*Kurchatov genomics center ICG SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Novosibirsk State University, Novosibirsk, Russia*

# SYMPOSIUM

## Bioinformatics and systems biology of plants

Oral reports

8 July, Wednesday

Small Conference Hall

Morning session. Bioinformatics and systems biology of plants

Chair: *Elena Salina, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia;*  
*Victoria Mironova, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia; NSU, Novosibirsk, Russia*

9.00 – 9.30

### PLENARY REPORT

**Human Genome, Anesthesiology and Critical Care. 20 Years Later**  
Vladimir Zelman,

- 9:40 – 10:00**      **Meta – analysis of transcriptome data clarified hormonal regulation of cold stress response in *Arabidopsis thaliana* L.**  
Nadezda Omelyanchuk<sup>1</sup>, Yana Sizentsova<sup>1</sup>, Victoria Mironova<sup>1,2</sup>  
<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*
- 10:00 – 10:20**      **Transcriptomic mechanisms of *Solanum tuberosum* defensive response to golden potato nematode infestation**  
Alexey Kochetov<sup>1</sup>, Kseniya Strygina<sup>2</sup>, Elena Khlestkina<sup>2</sup>, Egorova Anastasiya<sup>1</sup>, Dmitry Afonnikov<sup>1</sup>, Sophia Gerasimova<sup>1</sup>, Anastasiya Glagoleva<sup>1</sup>, Nickolay Shmakov<sup>1</sup>  
<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
<sup>2</sup>*VIR, St. Petersburg, Russia*
- 10:20 – 10:40**      **EIN3 binding site architecture guides transcriptional response to ethylene in *Arabidopsis***  
Vladislav Dolgikh<sup>1</sup>, Victor Levitsky<sup>1</sup>, Elena Zemlyanskaya<sup>1,2</sup>, Dmitry Oshchepkov<sup>1</sup>  
<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
<sup>2</sup>*NSU, Novosibirsk, Russia*
- 10:40 – 11:00**      **Features of the organization of bread wheat 5BS chromosome region carrying the leaf rust resistance gene *Lr52***  
Maria Braçina<sup>1</sup>, Dmitriy Afonnikov<sup>1,2</sup>, Elena Salina<sup>1</sup>  
<sup>1</sup>*Kurchatov Genomic Center, ICG SB RAS, Novosibirsk, Russia*  
<sup>2</sup>*NSU, Novosibirsk, Russia*
- 11:00 – 11:20**      **Symmetry and Asymmetry in Bacterial and Organellae Genomes**  
Michael Sadovsky, Maria Senashova  
*ICM SB RAS, Krasnoyarsk, Russia*
- 11:20 – 11:40**      **Coffee break**
- 11:40 – 12:10**      **Spikes morphometric characteristics analysis of five species of wheat**  
Evgeniy Komyshev<sup>1</sup>, Yuliya Kruchinina<sup>1</sup>, Mikhail Genaev<sup>1,2</sup>, Vasiliy Koval<sup>1</sup>, Dmitry Afonnikov<sup>1,2</sup>, Nikolay Goncharov<sup>3</sup>  
<sup>1</sup>*Kurchatov Genomic Center, ICG SB RAS, Novosibirsk, Russia*  
<sup>2</sup>*NSU, Novosibirsk, Russia*  
<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**12:10 – 12:30**      **Keeping the gate closed: WOX5 supports the balance between the proximal and distal root meristems via auxin biosynthesis in *Arabidopsis thaliana* L.**

Maria Savina<sup>1</sup>, Nadezda Omelyanchuk<sup>1</sup>, Taras Pasternak<sup>2</sup>, Victoria Mironova<sup>1,3</sup>, Viktoriya Lavrekha<sup>1</sup>

<sup>1</sup>ICG SB RAS, Novosibirsk, Russia

<sup>2</sup>Institute of Biology II/Molecular Plant Physiology University of Freiburg, Freiburg, Germany

<sup>3</sup>Novosibirsk State University

**12:30 – 12:50**      **Genetic regulation of wheat inflorescence development**

Oxana B. Dobrovolskaya<sup>1,2</sup>, Alina E. Dresvyannikova<sup>2</sup>, Petr Martinik<sup>3</sup>

<sup>1</sup>VNIIKR, Moscow region, Ramenskoe district, Bykovo, Russia

<sup>2</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>3</sup>Agrotest Fyto, Ltd, Kroměříž, Czech Republic

**12:50 – 13:10**      **Estimation of a joint distribution for several phenotypic traits in breeding or ancient populations**

Anna A. Igolkina<sup>1</sup>, Sergey Nuzhdin<sup>1,2</sup>, Maria G. Samsonova<sup>1</sup>

<sup>1</sup>SPbPU, St.Petersburg, Russia

<sup>2</sup>UCS, Los Angeles, USA

**13:10 – 14:30**      **Lunch**

**Evening session. Bioinformatics and systems biology of plants**

**Chairs: Elena Salina, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia;  
Elena Zemlyanskaya, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia; NSU, Novosibirsk, Russia**

**14.30 – 15.00**

**PLENARY REPORT**

**Understanding the evolution of complex regions in bird genomes**

Alexander Sang-Jae Suh,

Department of Evolutionary Biology, Uppsala University, Uppsala, Sweden

**15:00 – 15:30**

**Amyloidogenic properties of the beta – barrel proteins and their involvement in storage of nutrients in plant seeds and bacteria virulence**

Nizhnikov Anton<sup>1,2</sup>

<sup>1</sup>ARRIAM, St. Petersburg, Russia, St. Petersburg State University

<sup>2</sup>St. Petersburg State University, St. Petersburg, Russia

**15:30 – 15:50**

**Simulation climatic model for time to flowering in wild chickpea**

Andrey Ageev<sup>1</sup>, Abdullah Kahraman<sup>2</sup>, Sergey Nuzhdin<sup>1,3</sup>, Jens Berger<sup>4</sup>, Abdulkadir Aydogan<sup>5</sup>, Maria Samsonova<sup>1</sup>, Eric Bishop – von Wettberg<sup>6</sup>, Douglas Cook<sup>7</sup>, Konstantin Kozlov<sup>1</sup>

<sup>1</sup>SPbPU, St.Petersburg, Russia

<sup>2</sup>Harran University, Sanliurfa, Turkey

<sup>3</sup>USC, LA, CA, USA

<sup>4</sup>CSIRO, WA, Australia

<sup>5</sup>CRIFC, Ankara, Turkey

<sup>6</sup>UVM, VT, USA

<sup>7</sup>UC Davis, CA, USA

**15:50 – 16:10**

**MtWOX9 – 1 gene as somatic embryogenesis stimulator. Search of targets**

Varvara Tvorogova, Ksenia Kuznetsova, Elizaveta Krasnoperova, Elina Potsenkovskaya, Andrei Kudriashov, Ludmila Lutova  
SPSU, St.Petersburg, Russia

**16:10– 17:10**

**Coffee break**

**17:10 – 17:30**

**Identifying novel elements and regulators in auxin – dependent gene expression**

Daria Novikova<sup>1</sup>, Dolf Weijer<sup>2</sup>, Nadezda Omelyanchuk<sup>1</sup>, Victoria Mironova<sup>1</sup>

<sup>1</sup>Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia

<sup>2</sup>Wageningen University and Research, Wageningen, Netherlands

**17:30 – 17:50**

**Targeted mutagenesis of the *HvMyc2* and *HvAnt2* genes in *Hordeum vulgare* L.**

Anastasiya Egorova<sup>1,2,3</sup>, Christian Hertig<sup>3</sup>, Alexander Vikhorev<sup>1,2</sup>, Ksenia Strygina<sup>4</sup>, Iris Koeppel<sup>3</sup>, Sophia Gerasimova<sup>1,2</sup>, Elena Khlestkina<sup>4</sup>, Olesya Shoeva<sup>1</sup>, Stefan Hiekel<sup>3</sup>, Jochen Kumlehn<sup>3</sup>

<sup>1</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>2</sup>NSU, Novosibirsk, Russia

<sup>3</sup>IPK, Gatersleben, Germany

<sup>4</sup>VIR, St.Petersburg, Russia

**17:50 – 18:10**

**The meta – analysis of transcriptomes of Arabidopsis thaliana transgenic plants with altered expression of dual – targeting RNA – polymerase RPOTmp**

Igor Gorbenko, Vadim Belkov, Vladislav Tarasenko, Yuri Konstantinov, Alexander Katyshev, Milana Koulintchenko  
SIPPB SB RAS, Irkutsk, Russia

18:10 – 18:30

**Transcripts Specifically Expressed During Secondary Vascular Development in *Arabidopsis thaliana* L.**

Nadezda Omelyanchuk<sup>1,2</sup>, Dmitry Oshchepkov<sup>1</sup>, Evgeniya Pukhovaya<sup>1,2</sup>, Victoria Mironova<sup>1</sup>

<sup>1</sup>*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

18:30 – 18:50

**Exploring Interaction Between Metabolic Pathways Involved In Pigmentation Of Barley Spike**

Anastasiia Glagoleva<sup>1</sup>, Nikolay A. Shmakov<sup>1</sup>, Aleksandr V. Vikhorev<sup>1,2</sup>, Sergei R. Mursalimov<sup>1</sup>, Natalia V. Gracheva<sup>3</sup>, Tatjana V.

Kukoeva<sup>1</sup>, Olesya Yu. Shoeva<sup>1</sup>, Elena K. Khlestkina<sup>1,4</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*VSTU, Volgograd, Russia*

<sup>4</sup>*VIR, Saint – Petersburg, Russia*

18:50 – 19:00

**Development of DNA markers for identification of a quarantine weed, silverleaf nightshade (*Solanum elaeagnifolium* Cav.), based on chloroplast intergenic spacers**

Ekaterina Volodina<sup>1</sup>, Y.Y. Kulakova<sup>1</sup>, O. B. Dobrovolskaya<sup>1,2</sup>, M.S. Anisimenko<sup>1</sup>

<sup>1</sup>*VNIKR, Bykovo, Russia*

<sup>2</sup>*ICG SB RAS, Novosibirsk, Russia*

**Poster session**

**Effects of anthocyanin – rich grain diet on growth and metastasis of Lewis lung carcinoma in mice**

Michael V. Tenditnik<sup>1</sup>, Nelly A. Popova<sup>2</sup>, Maria A. Tikhonova<sup>1</sup>, Tamara G. Amstislavskaya<sup>1</sup>, Ekaterina A. Litvinova<sup>1</sup>, Elena K. Khlestkina<sup>3,2</sup>

<sup>1</sup>*Scientific Research Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

<sup>2</sup>*Federal Research Center “Institute of Cytology and Genetics”, Novosibirsk, Russia*

*N.I. Vavilov All – Russian Research Institute of Plant Genetic Resources, St. Petersburg, Russia*

**Molecular markers based on SNPs in *FAD3* genes for determination of linolenic acid content in flax seed**

Liubov Povkhova<sup>1,2</sup>, Elena Pushkova<sup>1</sup>, Alexey Dmitriev<sup>1</sup>, Parfe Kezimana<sup>1,3</sup>, Roman Novakovskiy<sup>1</sup>, Nataliya Melnikova<sup>1</sup>, Tatiana Rozhmina<sup>1,4</sup>, George Krasnov<sup>1</sup>

<sup>1</sup>*Engelhardt Institute of Molecular Biology, RAS, Moscow, Russia;*

<sup>2</sup>*Moscow Institute of Physics and Technology, Dolgoprudny, Russia*

<sup>3</sup>*RUDN University, Moscow, Russia*

<sup>4</sup>*Federal Research Center for Bast Fiber Crops, Torzhok, Russia*

**Flowering patterns of herbaceous multi – flowered monocarpic shoots of *Campanula sarmatica***

Fomin Eduard Fomin

### **Wheat and maize miRNAs are potential regulators of human genes expression**

Aizhan Kazievna Rakhmetullina, Anatoliy Timofeevich Ivashchenko, Anna Yurevna Pyrkova

*AI – Farabi Kazakh National University, Almaty, Kazakhstan*

### **The characteristics of interaction of miRNA with mRNA of C2H2, ERF and GRAS transcription factors of arabidopsis, rice and maize**

Aizhan Kazievna Rakhmetullina, Svetlana Kazbekovna Turasheva, Anna Yurevna Pyrkova

*AI – Farabi Kazakh National University, Almaty, Kazakhstan*

### **Genome – wide Prediction of Transcription Start Site in Four Conifer Species**

Eugeniia I. Bondar<sup>1,2</sup>, Vadim V. Sharov<sup>1,2</sup>, Dmitry A. Kuzmin<sup>1</sup>, Tatiana V. Tatarinova<sup>1,3,4,5</sup>, Konstantin V. Krutovsky<sup>1,6,4,7</sup>

<sup>1</sup>*Siberian Federal University, Krasnoyarsk, Russia*

<sup>2</sup>*FRC KSC SB RAS, Krasnoyarsk, Russia*

<sup>3</sup>*University of La Verne, La Verne, USA*

<sup>4</sup>*Vavilov Institute of General Genetics, Moscow, Russia*

<sup>5</sup>*Bioinformatics Center of IITP RAS, Moscow, Russia*

<sup>6</sup>*Georg – August University of Göttingen Göttingen, Germany*

<sup>7</sup>*Texas A&M University, College Station, TX, USA*

### **Molecular genetic analysis of alloplasmic recombinant lines (*Triticum dicoccum*) – *Triticum aestivum***

Andrey Borisovich Shcherban<sup>1</sup>, Roman Nikolaevich Perfil'ev<sup>2</sup>, Elena Artemovna Salina<sup>1</sup>

<sup>1</sup>*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State Agrarian University*

### **Genome – wide analysis of highly expressed plant retrotransposons**

Murad Omarov, Pavel Merkulov, Sofia Gvaramia, Liza Kolganova, Ilya Kirov

*All – Russia Research Institute of Agriculture biotechnology, Moscow, Russia*

### **The prospects for the study of the avirulence genes characteristic for the West Siberian population of wheat stem rust *Puccinia graminis* f. sp. *Tritici***

Vasiliy Kelbin, Ekaterina Sergeeva, Elena Salina, Ekaterina Skolotneva

*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

### **Retrotransposons of *Arabidopsis thaliana* expressed in wild – type plants**

Sofya Gvaramiya, Murad Omarov, Ilya Kirov

*All – Russia Research Institute of Agricultural Biotechnology, Moscow, Russia*

### **Identification of an AP2/ERF Transcription Factor Controlling the Synthesis of Barley Epicuticular Wax**

Ekaterina Kolosovskaya<sup>1</sup>, Christian Hertig<sup>2</sup>, Dmitriy Domrachev<sup>3</sup>, Alexey Kochetov<sup>1</sup>, Sophia Gerasimova<sup>4</sup>, Sergey Morozov<sup>3</sup>, Vikhorev Alexander<sup>1,4</sup>, Jochen Kumlehn<sup>2</sup>, Anna Korotkova<sup>1</sup>, Elena Chernyak<sup>2</sup>, Nikolay Shmakov<sup>1</sup>, Elena Khlestkina<sup>1,4,5</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Leibniz Institute of Plant Genetics and Crop Plant Research (IPK), Gatersleben, Germany*

<sup>3</sup>*Novosibirsk Institute of Organic Chemistry, SB RAS, Novosibirsk, Russia*

<sup>4</sup>*NSU, Novosibirsk, Russia*

<sup>5</sup>*Vavilov Institute of Plant Genetic Resources (VIR), Saint Petersburg, Russia*

### **Genomic analysis of Vavilov's historic chickpea landraces using GWAS, AMMI and GGE biplot analyses**

Alena Sokolkova<sup>1</sup>, Noelia Carrasquilla – Garcia<sup>2</sup>, Douglas R. Cook<sup>2</sup>, Sergey V. Bulyntsev<sup>3</sup>, Eric von Wettberg<sup>4</sup>, Sergey V. Nuzhdin<sup>5</sup>, Peter L. Chang<sup>5</sup>, Margarita A. Vishnyakova<sup>3</sup>, Maria G. Samsonova<sup>6</sup>

<sup>1</sup>*Peter the Great St. Petersburg Polytechnic University, St. Petersburg, Russia*

<sup>2</sup>*University of California Davis, Department of Plant Pathology, Davis, CA 95616 USA*

<sup>3</sup>*Federal Research Centre All – Russian N.I. Vavilov Institute of Plant Genetic Resources (VIR), St. Petersburg, Russia*

<sup>4</sup>*University of Vermont, Department of Plant and Soil Science, Burlington, VT 05405, USA*

<sup>5</sup>*University of Southern California, Program in Molecular and Computational Biology, Dornsife College of Letters Arts & Sciences, Los Angeles, USA*

<sup>6</sup>*Peter the Great St. Petersburg Polytechnic University, Department of Applied Mathematics, St. Petersburg, Russia*

### **Analysis of agronomic traits of mungbean (*Vigna radiata*) accessions from the World Vegetable Gene Bank (Taiwan)**

Alena Sokolkova<sup>1</sup>, Margarita A. Vishnyakova<sup>2</sup>, Chau – Ti Ting<sup>3</sup>, Marina Burlyaeva<sup>2</sup>, Roland Schafleitner<sup>4</sup>, Sergey V. Nuzhdin<sup>5</sup>, Eric von Wettberg<sup>6</sup>, Tatjana Valiannikova<sup>7</sup>, Cheng – Rwei Lee<sup>3</sup>, Maria G. Samsonova<sup>8</sup>

<sup>1</sup>*Peter the Great St. Petersburg Polytechnic University, St. Petersburg, Russia*

<sup>2</sup>*Federal Research Centre All – Russian N.I. Vavilov Institute of Plant Genetic Resources (VIR), St. Petersburg, Russia*

<sup>3</sup>*National Taiwan University, Taipei 106, Taiwan*

<sup>4</sup>*World Vegetable Center, Shanhua, Tainan 74199, Taiwan*

<sup>5</sup>*University of Southern California, Program in Molecular and Computational Biology, Dornsife College of Letters Arts & Sciences, Los Angeles, USA*

<sup>6</sup>*University of Vermont, Department of Plant and Soil Science, Burlington, VT 05405, USA*

<sup>7</sup>*Kuban Branch of Federal Research Centre All – Russian N.I. Vavilov Institute of Plant Genetic Resources (VIR), Krasnodar region, Russia*

<sup>8</sup>*Peter the Great St. Petersburg Polytechnic University, Department of Applied Mathematics, St. Petersburg, Russia*

### **STUDY OF THE ROOT TRANSCRIPTOME OF BREAD WHEAT USING HIGH – THROUGHPUT RNA SEQUENCING (RNA – SEQ)**

Alexandr Vikhorev<sup>1,2</sup>, Elena Khlestkina<sup>2,3</sup>, Nikolay Shmakov<sup>2</sup>, Olesya Shoeva<sup>2</sup>, Anastasia Glagoleva<sup>2</sup>

<sup>1</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

<sup>3</sup>*All – Russian Institute of Plant Recources, Saint – Petersburg, Russia*

### **Btr1 genes and the evolution of wheat and *Aegilops* species**

Valeriya Vavilova, Irina Konopatskaia, Nikolay P. Goncharov, Alexandr Blinov

<sup>2</sup>*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

### **Regulation of Transcription Activity of MAKR4 in *Arabidopsis thaliana* L.**

Anastasia Korosteleva<sup>1</sup>, Daria Novikova<sup>2</sup>, Victoria Mironova<sup>2</sup>

<sup>1</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

### **Analysis of repeatomes in Cannabaceae family**

Julia Bocharkina<sup>1,2</sup>, Olga Razumova<sup>1</sup>, Gennady Karlov<sup>1</sup>

<sup>1</sup>*All – Russia Research Institute of Agricultural Biotechnology, Moscow, Russia*

<sup>2</sup>*Skolkovo Institute of science and technology, Moscow, Russia*

### **Comparative genomic analysis of male and female poplars**

Elena Pushkova, Nadezhda Bolsheva, George Krasnov, Nataliya Melnikova, Roman Novakovskiy, Alexey Dmitriev

*Engelhardt Institute of Molecular Biology, RAS, Moscow, Russia*

### **Comparative analysis of flax (*Linum usitatissimum* L.) genomes and transcriptomes**

Elena Pushkova<sup>1</sup>, Liubov Povkhova<sup>1,2</sup>, Tatiana Rozhmina<sup>1,3</sup>, George Krasnov<sup>1</sup>, Artemy Beniaminov<sup>1</sup>, Alexey Dmitriev<sup>1</sup>, Roman Novakovskiy<sup>1</sup>, Nadezhda Bolsheva<sup>1</sup>, Nataliya Melnikova<sup>1</sup>

<sup>1</sup>*Engelhardt Institute of Molecular Biology, RAS, Moscow, Russia*

<sup>2</sup>*Moscow Institute of Physics and Technology, Dolgoprudny, Russia*

<sup>3</sup>*Federal Research Center for Bast Fiber Crops, Torzhok, Russia*

### **Complete sequencing of barley organellar genomes: new data for intraspecific differentiation**

Yermakovich (Makarevich) Anna, Liaudanski Aleh, Siniauskaya Maryna, Davydenko Oleg, Halayenka Innesa *IGS NAS of Belarus, Minsk, Belarus*

### **SeedCounter – mobile application for high throughput grain phenotyping**

Mikhail Genaev<sup>1,2</sup>, Komyshev Evgeny<sup>1</sup>, Dmitry Afonnikov<sup>1,2</sup>

<sup>1</sup>*ICG SB RAS, Kurchatov Genomic Center, Novosibirsk, Russia*

<sup>2</sup>*NSU, Novosibirsk, Russia*

### **Application of neural networks to image recognition of wheat rust diseases**

Mikhail Genaev<sup>1,2</sup>, Skolotneva Ekaterina<sup>1</sup>, Dmitry Afonnikov<sup>1,2</sup>

<sup>1</sup>*ICG SB RAS, Kurchatov Genomic Center, Novosibirsk, Russia*

<sup>2</sup>*NSU, Novosibirsk, Russia*

### **Detection and investigation of genes with circadian expression pattern in common wheat**

Antonina Kiseleva, Maria Bragina, Elena Salina  
*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

**Identification of genetic factors responsible for symbiotic effectiveness in pea (*Pisum sativum* L.)**

Aleksandr Zhernakov<sup>1</sup>, Igor Tikhonovich<sup>1,2</sup>, Oksana Shtark<sup>1</sup>, Vladimir Zhukov<sup>1</sup>, Olga Kulaeva<sup>1</sup>

<sup>1</sup>*ARRIAM, St. Petersburg, Russia*

<sup>2</sup>*Saint – Petersburg State University, Saint – Petersburg, Russia*

**New insight on diversity of the Nikita Botanical Gardens plant collections from advanced NGS**

Irina Mitrofanova, Svetlana Chelombit, Olga Krivenko, Valentina Brailko, Olga Kuleshova, Olga Mitrofanova

*NBG – NSC RAS, Yalta, Russia*

## SECTION Structural Computational Biology

**Oral reports**

9 July, Thursday

**Computer Class**

**Chairs:** Vladimir Poroikov, *Institute of Biomedical Chemistry, Moscow, Russia*

Vladimir Ivanisenko, *Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia; Novosibirsk State University, Novosibirsk, Russia*

14.30 – 15.00

**PLENARY REPORT**

**Brain Asymmetries in the Cognitive Processing**

Boris M. Velichkovsky

*Corresponding Member of the Russian Academy of Science, Member of Academia Europaea*

15:00 – 15:30	<p><b>Keynote report</b>  <b>Computer – aided approaches to discovery of novel pharmaceutical agents for COVID – 19 therapy</b>  <u>Vladimir Poroikov</u>,  <i>Institute of Biomedical Chemistry, Moscow, Russia</i></p>
15:30 – 15:50	<p><b>Computer tools for modelling and prediction of natural RNA structure: a case study of miRNAs and group II introns</b>  <u>Igor Titov</u>  <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i></p>
15:50 – 16:10	<p><b>Computer analysis of aminoacid residue patterns in protein 3d structure similar to functional sites.</b>  <u>V.A. Ivanisenko</u>, N.V. Ivanisenko, N.A. Kolchanov  <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i></p>
16:10 – 16:30	<p><b>Learning the changes of barnase mutants thermostability from structural fluctuations obtained using anisotropic network modeling</b>  <u>Nikolay Alemasov</u>, Nikita Ivanisenko, Vladimir Ivanisenko  <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i></p>
16:30 – 16:40	<p><b>Coffee – break</b></p>
16:40 – 17:00	<p><b>Mustguseal: versatile bioinformatic platform for knowledge – based protein design and modulation</b>  <u>Dmitry Suplatov</u><sup>1,2</sup>, Yana Sharapova<sup>1,2</sup>, Vytas Švedas<sup>1,2</sup>  <sup>1</sup><i>Belozersky Institute of Physicochemical Biology, Moscow, Russia</i>  <sup>2</sup><i>Lomonosov Moscow State University, Moscow, Russia</i></p>
17:00 – 17:20	<p><b>Model for Stacking Monomers in Filamentous Actin</b>  <u>Anna Glyakina</u><sup>1,2</sup>, Alexey Surin<sup>2,3</sup>, Oxana Galzitskaya<sup>2</sup>  <sup>1</sup><i>IMPB RAS, Pushchino, Russia</i>  <sup>2</sup><i>Institute of Protein Research RAS, Pushchino, Russia</i>  <sup>3</sup><i>Pushchino Branch, Shemyakin–Ovchinnikov Institute of Bioorganic Chemistry, RAS, Pushchino, Russia</i></p>
17:20 – 17:40	<p><b>Parallel Bias Metadynamics and Sketch – map Dimensionality Reduction as Powerful Tools to Explore Free Energy Landscapes of Intrinsically Disordered Peptides</b>  <u>Olga Rogacheva</u><sup>1,2</sup>, Omar Valsson<sup>3</sup>, Olga Shamova<sup>2</sup>, Andrey Badanin<sup>1</sup>  <sup>1</sup><i>SPbU, St. Petersburg, Russia</i>  <sup>2</sup><i>FSBRI “IEM”, St. Petersburg, Russia</i>  <sup>3</sup><i>MPIP, Mainz, Germany</i></p>

17:40 – 18:00	<b>An effective molecular blockers of ion channel of M2 protein as anti – influenza A drug</b> Yury Nikolaevich Vorobjev <i>Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia</i>
18:00 – 18:10	<b>Coffee – break</b>
18:10 – 18:30	<b>Extraction of spectral series of ions from mass spectra of peptides by methods of integral transforms and machine learning</b> <u>Eduard Fomin</u> , Nikolay Alemasov, Dmitriy Afonnikov <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i>
18:30 – 18:50	<b>Modeling of single – molecule FRET – experiments on protein folding: From coarse – grained to all – atom simulations</b> <u>Vladimir A. Andryushchenko</u> <sup>1,2</sup> , Sergei F. Chekmarev <sup>1</sup> <sup>1</sup> <i>IT SB RAS, Novosibirsk, Russia</i> <sup>2</sup> <i>NSU, Novosibirsk, Russia</i>
18:50 – 19:10	<b>3D agent – based modeling of some aspects of the interaction between microtubules and microfilaments in cell</b> <u>Marat Sabirov</u> , Alexander Spirov <i>I.M. Sechenov Institute of Evolutionary Physiology and Biochemistry Russian Academy of Sciences Saint – Petersburg, Russia</i>
19:10 – 19:30	<b>Advanced laser technologies for targeted nuclear nanomedicine</b> Irina Zavestovskaya <i>National Research Nuclear University MEPhI, Moscow, Russia</i> <i>The Lebedev Physical Institute, Moscow, Russia</i>

## Poster session

### Searching for Alternatively Splicing Group II Introns

Nikolay Kobalo<sup>1</sup>, Denis Vorobyev<sup>2</sup>, Igor Titov<sup>3</sup>, Alexander Kulikov<sup>1</sup>

<sup>1</sup>*The Institute of Computational Mathematics and Mathematical Geophysics, Novosibirsk, Russia*

<sup>2</sup>*Gustave Roussy Cancer Center, Villejuif, France*

<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### The properties of the C – terminal domain of HlyIICTD suggest that *B. cereus* HlyII is a representative potential member of trimeric autotransporter adhesins among gram – positive bacteria.

Siunov A.V. <sup>a</sup>, Nagel A. S. <sup>a</sup>, Andreeva – Kovalevskaya Z. I. <sup>a</sup>, Zamyatina A.V. <sup>b, c</sup>, Rudenko N.V. <sup>b, c</sup>, Karatovskaya A.P. <sup>c</sup>, Borisova M. P. <sup>d</sup>, Brovko F.A. <sup>b, c</sup>, Salyamov V. I. <sup>a</sup>, A.S. Solonin <sup>a, b</sup>

*a G. K. Skryabin Institute of Biochemistry and physiology of microorganisms RAS 5 Prospekt Nauki, Pushchino, Moscow Region 142290, Russia*

*b Pushchino State Institute of Natural Sciences, 3 Prospekt Nauki, Pushchino, Moscow Region 142290, Russia*

*c Pushchino Branch, Shemyakin–Ovchinnikov Institute of Bioorganic Chemistry, 6 Prospekt Nauki, Pushchino, Moscow Region 142290, Russia*  
*d Institute of Theoretical and Experimental Biophysics, Russian Academy of Sciences, Pushchino, Moscow region, 142290, Russia*

### **Competition and collaboration in the miRNA science field**

*Artemiy Firsov<sup>1</sup>, Igor Titov<sup>2</sup>*

<sup>1</sup>*Computer Science and Computer Engineering, Institute of Informarics Systems, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **Errors in miRNA Recognition**

*Pavel Vorozheykin<sup>1</sup>, Igor Titov<sup>1,2</sup>*

<sup>1</sup>*NSU, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **An architecture – independent algorithm for microRNA target prediction**

*Natalya Fokina, Alexander Grinev*

*Moscow State Medical University Moscow, Russia*

### **Nuclear envelope rupture in *Drosophila* D11 cells inhibit mitosis**

*Snezhanna Sergeevna Saydakova<sup>1,2</sup>, Gera Alekseevna Pavlova<sup>3</sup>, Ksenia Nikolaevna Morozova<sup>1</sup>, Elena Vladimirovna Kiseleva<sup>1</sup>*

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*NSU, Novosibirsk, Russia*

<sup>3</sup>*IMCB SB RAS Novosibirsk, Russia*

### **Consideration of pathogenicity of nsSNVs in CDKN2A gene, as a new tumor marker for leukemia, using bioinformatics methods**

*Farzaneh Ghasemi<sup>1</sup>, Mehri Khatami<sup>1</sup>, Mohammad Mehdi Heidari<sup>1</sup>, Yuriy L. Orlov<sup>2,3</sup>*

<sup>1</sup>*Yazd University, Yazd, Iran*

<sup>2</sup>*I.M. Sechenov First Moscow State Medical University, Moscow, Russia*

<sup>3</sup>*Novosibirsk State University, Novosibirsk, Russia*

### **easyAmber: a step away from inefficient “static” approaches towards a deeper understanding of protein dynamics**

*Dmitry Suplatov<sup>1,2</sup>, Yana Sharapova<sup>1,2</sup>, Vytas Švedas<sup>1,2</sup>*

<sup>1</sup>*Belozersky Institute of Physicochemical Biology, Moscow, Russia*

<sup>2</sup>*Lomonosov Moscow State University, Moscow, Russia*

### **Beta – Bends As An Example Of Conformationally Predetermined Segments Of Protein. Conditions Of Stabilization Of The Structure And Role Of Context**

*Anastasia A. Anashkina<sup>1</sup>, Vladimir O. Chekhov<sup>1</sup>, Ivan Yu. Torshin<sup>2</sup>, Leonid A. Uroshlev<sup>3</sup>, Natalia G. Esipova<sup>1</sup>, Vladimir G. Tumanyan<sup>1</sup>*

<sup>1</sup>*EIMB RAS, Moscow, Russia*

<sup>2</sup>*FIC IU RAS, Moscow, Russia*

<sup>3</sup>*IGG RAS, Moscow, Russia*

**Modelling of Nef Interaction with ABCA1 Revealed Potential Binding Sites For Inhibitor Compounds**

*Anastasia A. Anashkina, Yaroslav V. Tkachev, Alexei A. Adzhubei  
EIMB RAS, Moscow, Russia*

**Computer reconstruction of the ecological structure of intestinal microbiota communities based on high – throughput sequencing data**

*Andrew Kopochev<sup>1</sup>, Alexandra Klimenko<sup>1</sup>, S.A. Lashin<sup>1,2</sup>*

*<sup>1</sup>Kurchatov Genomics Center, Institute of Cytology and Genetics, ICG SB RAS, Novosibirsk, Russia*

*<sup>2</sup>Novosibirsk State University, NSU, Novosibirsk, Russia*

**Interpretation of the features of a linear regression model for predicting the survival time of the amyotrophic lateral sclerosis patients with mutated SOD1**

*Nikolay Alemasov<sup>1</sup>, Shcherbakov Alexandr<sup>2</sup>, Vladimir Timofeev<sup>2</sup>, Vladimir Ivanisenko<sup>1</sup>*

*<sup>1</sup> Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

*<sup>2</sup>Novosibirsk State Technical University Novosibirsk, Russia*

# SECTION

## Systems Biology of Aging

Oral reports

### 9 July, Thursday

Small Conference Hall

Chairs: Elena Pasyukova, *IMG RAS, Moscow, Russia*  
Vyacheslav Mordvinov, *Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
Alexander Khokhlov, *Lomonosov Moscow State University, Moscow, Russia*

14.30 – 15.00

#### **PLENARY REPORT**

##### **Brain Asymmetries in the Cognitive Processing**

Boris M. Velichkovsky

*Corresponding Member of the Russian Academy of Science, Member of Academia Europaea*

15:00 – 15:05

Welcoming address by the President of the Gerontological Society of the Russian Academy of Sciences Vladimir Anisimov, St. Petersburg, Russia

15:05 – 15:30

#### **Neuronal Transcription Factors in Lifespan Control**

Alexander Symonenko, Natalia Roshina, Anna Kremetsova, Elena Pasyukova

*IMG RAS, Moscow, Russia*

15:30 – 15:50

#### **Evolution of Proteins Involved in Response to ROS**

Vassily Lyubetsky<sup>1</sup>, Gregory Shilovsky<sup>1,2</sup>, Alexandr Seliverstov<sup>1</sup>, Oleg Zverkov<sup>1</sup>, Lev Rubanov<sup>1</sup>

<sup>1</sup>*Institute for information transmission problems, RAS, Moscow, Russia*

<sup>2</sup>*Lomonosov Moscow State University, Moscow, Russia*

15:50 – 16:10

#### **Gerontology and Scientometrics ("Citogerontology")**

Alexander Khokhlov

*Lomonosov Moscow State University, Moscow, Russia*

- 16:10 – 16:30**      **Cholinergic Deficit in Olfactory Bulbectomized Animals as a Model of Neurodegenerative Diseases**  
Mikhail Stepanichev, Olga Nedogreeva, Natalia Lazareva, Anna Manolova, Natalia Gulyaeva  
*IHNA&NPh RAS, Moscow, Russia*
- 16:30 – 16:50**      **Cellular senescence in age – related macular degeneration: impact of changes in autophagy and neurotrophic supplementation**  
Oyuna Kozhevnikova, Darya Telegina, Mikhail Tyumentsev, Nataliya Kolosova  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*
- 16:50 – 17:10**      **Coffee break**
- 17:10 – 17:40**      **Free Radical Theory of Aging: from Chemical Physics to Systems Theory of Reliability**  
*Vitaly Koltover*  
*Institute of Problems of Chemical Physics RAS, Chernogolovka, Russia*
- 17:40 – 18:00**      **Cluster analysis of age – related trends of the expression of metabolically relevant genes in humans PBMCs**  
Aleksy Alekseev  
*Lomonosov Moscow State University, Moscow, Russia*
- 18:00 – 18:20**      **Cellular response to UVA – B light depends on cellular age and chromatin structure**  
Bela Vasileva<sup>1</sup>, Dessislava Staneva<sup>1</sup>, Plamen Zagorchev<sup>2</sup>, Natalia Krasteva<sup>3</sup>, George Miloshev<sup>1</sup>, Milena Georgieva<sup>1</sup>  
<sup>1</sup>*Institute of Molecular Biology “Acad. R. Tsanev” Bulgarian Academy of Sciences, Sofia, Bulgaria*  
<sup>2</sup>*Medical University – Plovdiv, Plovdiv, Bulgaria*  
<sup>3</sup>*Institute of Biophysics and Biomedical Engineering, Bulgarian Academy of Sciences, Sofia, Bulgaria*
- 18:20 – 18:40**      **Is there a fecundity/longevity trade – off under heat stress?**  
Nataly Gruntenko, Evgenia Karpova, Elena Burdina, Natalya Adonyeva, Petr Menshanov, Inga Rauschenbach  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*
- 18:40 – 19:00**      **Delay of early postnatal development as a risk factor for accelerated aging and Alzheimer’s disease**  
Ekaterina Rudnitskaya, Tatiana Kozlova, Alena Burnyasheva, Natalia Stefanova, Nataliya Kolosova  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*
- 19:00 – 19:20**      **Serum Polypeptide Alpha – Fetoprotein (AFP) as a Possible Powerful Geroprotector**  
Alexander Khalyavkin<sup>1,2</sup>, Vyacheslav Krut’ko<sup>2,3</sup>, Vitaly Dontsov<sup>2</sup>  
<sup>1</sup>*Institute of Biochemical Physics of RAS, Moscow, Russia*  
<sup>2</sup>*Federal Research Center “Computer Science and Control” of RAS Moscow, Russia*  
<sup>3</sup>*Sechenov First Moscow State Medical University Moscow, Russia*

**Possibility to use divergent tasks for baseline alpha rhythm modulation in older adults**

Evgeniya Privodnova, Nina Volf, Ekaterina Merculova, Victoriya Bilik  
*Scientific Research Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

**Spatial learning as activator of hippocampal neurogenesis during aging and development of Alzheimer's disease – like pathology**

Alena Burnyasheva, Tatiana Kozlova, Ekaterina Rudnitskaya, Natalia Stefanova  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**Effects of melatonin and SkQ1 long – term treatment during aging and development AMD – like retinopathy**

Darya Telegina, Oyuna Kozhevnikova, Anzhella Fursova  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**Age – related difference in use – dependent plasticity after divergent thinking session matches posterior – anterior shift in aging (PASA) model**

Evgeniya Privodnova, Nina Volf, Ekaterina Merculova, Dariya Bazovkina  
*Scientific Research Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

**Calorie Restriction in Gerontological Experiments on Cell Cultures**

Galina Morgunova, Alexander Khokhlov  
*Lomonosov Moscow State University, Moscow, Russia*

**Lymph nodes morphology as predictor natural and premature aging**

Olga Gorchakova<sup>1</sup>, Vladimir Gorchakov<sup>1,2</sup>, Georgy Demchenko<sup>3</sup>

<sup>1</sup>Research institute of a clinical and experimental lymphology – branch of Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>2</sup>Novosibirsk State University, Novosibirsk, Russia

<sup>3</sup>Institute of Physiology of Human and Animals of Committee of Science of the Ministry of Education and Science of the Republic of Kazakhstan, Almaty, Kazakhstan

**Nanobodies design for treatment of age – related diseases**

Mohammad Mehdi Heidari<sup>1</sup>, Yuriy Orlov<sup>2,3</sup>

<sup>1</sup>Yazd University, Yazd, Iran

<sup>2</sup>Institute of Digital Medicine I.M.Sechenov First Moscow State Medical University, Moscow, Russia

<sup>3</sup>Novosibirsk State University, Novosibirsk, Russia

**MAPK pathways and alphaB – crystallin phosphorylation in brain: a focus on aging and Alzheimer's disease**

Natalia Muraleva  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**Mitochondrial dysfunction and redox balance alterations in the development of AD – like pathology in OXYS rats**

Mikhail Tyumentsev<sup>1</sup>, Natalia Muraleva<sup>1</sup>, Yulia Polienko<sup>2</sup>, Artyom Gorodetsky<sup>2</sup>, Elena Bagryanskaya<sup>2</sup>  
*ICG SB RAS; <sup>2</sup>NIOCH SB RAS, Novosibirsk, RUSSIA*

**Search for single nucleotide polymorphisms (SNPs) associated with hypertension in the genome of senescence – accelerated OXYS rats**

Vasily Devyatkin, Natalia Muraleva, Olga Redina, Nataliya Kolosova  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**Way to longevity: role of antioxidant defense gene polymorphisms in successful adaptation**

Vera Erdman<sup>1</sup>, Timur Nasibullin<sup>1</sup>, Ilsa Tuktarova<sup>1</sup>, Ksenia Danilko<sup>1</sup>, Olga Mustafina<sup>1</sup>, Tatiana Viktorova<sup>1</sup>, Alisa Matua<sup>2</sup>  
<sup>1</sup>*IBG UFRC RAS, Ufa, Russia*  
<sup>2</sup>*SRI EPT ASA, Sukhum, Abkhazia*

# SYMPOSIUM

## Genomics, bioinformatics and evolution

Oral reports

9 July, Thursday

Big Conference Hall

Morning session. Genomics, bioinformatics and evolution

Chairs: Igor Rogozin, *National Institutes of Health, USA*  
Dmitry Afonnikov, *Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

9.00 – 9.30

### PLENARY REPORT

**Genetic Development of the Nociceptive System and the Epigenetics of the Experience of Pain**

Jack M. Berger

*Keck School of Medicine of USC, University of Southern California, Los Angeles, California, USA*

9:30 – 10:00

### Keynote report

**Stop codons within prokaryotic protein – coding genes: Indication of frequent read – through events**

*Igor B. Rogozin*

*NCBI, Bethesda, USA*

10:00 – 10:20

### Homologous series. Law or rule?

*Valentine Suslov, Mikhail Ponomarenko, Dmitry Rasskazov*

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

10:20 – 10:40

### Heat shock protein 90 as a long – term buffer of mutational burden

Valeria Timonina<sup>1</sup>, Evgenii Tretiakov<sup>2</sup>, Anastasia Sokol<sup>1</sup>, Dmitry Knorre<sup>3,4</sup>, Konstantin Gunbin<sup>5</sup>, Konstantin Popadin<sup>1,6</sup>

<sup>1</sup>*Immanuel Kant Baltic Federal University, Kaliningrad, Russia*

<sup>2</sup>*Medical University of Vienna, Vienna, Austria*

<sup>3</sup>*Belozersky Institute of Physico – Chemical Biology, Moscow, Russia*

<sup>4</sup>*Moscow State University, Moscow, Russia*

<sup>5</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>6</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*

10:40 – 11:00

**Polygenic Transmission Disequilibrium of slightly – deleterious variants in Down Syndrome trios**

Kseniia Sholokhova<sup>1</sup>, Viktor Shamansky<sup>1</sup>, Konstantin Gunbin<sup>1,2</sup>, Konstantin Popadin<sup>1,3</sup>

<sup>1</sup>*IKBFU, Kaliningrad, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*

11:00 – 11:20

**Tandem repeats are selfish elements which mark the level of hidden recombination in animal mitochondrial genomes**

Aleksandr Smirnov<sup>1</sup>, Konstantin Gunbin<sup>1,2</sup>, Alina A. Mikhailova<sup>1</sup>, Konstantin Popadin<sup>1,3</sup>, Valeria Lobanova<sup>1</sup>

<sup>1</sup>*Immanuel Kant Baltic Federal University, Kaliningrad, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*

11:20 – 11:40

**Coffee break**

11:40 – 12:00

**Analysis of the Associations between Missense Substitutions in the Human MT – ATP6 gene**

Maria Golubenko, Alexey Zarubin

*Research Institute of Medical Genetics, Tomsk NRMC, Tomsk, Russia*

12:00 – 12:20

**The genomes and mechanisms of adaptation to the cold climates in Russian native cattle breeds**

Laura Buggiotti<sup>1</sup>, Andrey Yurchenko<sup>2</sup>, Nikolay Yudin<sup>2</sup>, Denis M. Larkin<sup>1,2</sup>

<sup>1</sup>*Royal Veterinary College, University of London, London, UK*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

12:20 – 12:40

**Resequencing genomes of the Russian native Baikal and Tuva sheep breeds**

James Sweet – Jones<sup>1</sup>, Nikolay Yudin<sup>2</sup>, Denis M. Larkin<sup>1,2</sup>

<sup>1</sup>*Royal Veterinary College, University of London, London, UK*

<sup>2</sup>*Institute of Cytology and Genetics, Novosibirsk, Russia*

12:40 – 13:00

**The genetic component of the human embryonic selection: uncovering of the strength and main targets**

Sergey Oreshkov<sup>1</sup>, Evgenii Tretiakov<sup>2</sup>, Dmitrii Iliushchenko<sup>1</sup>, Elisaveta Zezyulya<sup>1</sup>, Konstantin Gunbin<sup>1,3</sup>, Konstantin Popadin<sup>1,4</sup>

<sup>1</sup>*Immanuel Kant Baltic Federal University, Kaliningrad, Russia*

<sup>2</sup>*Medical University of Vienna, Vienna, Austria*

<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>4</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*

13:00 – 13:20

**mtDNA mammalian evolution: mice walk with many little steps while elephants with a few big ones**

Dmitrii Iliushchenko<sup>1</sup>, Anastasia Sokol<sup>1</sup>, Konstantin Gunbin<sup>1,2</sup>, Konstantin Popadin<sup>1,3</sup>

<sup>1</sup>*Immanuel Kant Baltic Federal University, Kaliningrad, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*

13:20 – 13:35

**Laboratory of Genetic Analysis**

Akopyan Nare Akopovna, *Department of Biochemistry and Biotechnology, Himmed Company, Moscow, Russia*

Sponsor report – **Himmed Company**

Lunch

Evening session. Genomics, bioinformatics and evolution

Chairs: Vyacheslav Yurchenko, *University of Ostrava, Faculty of Science, Ostrava, Czech Republic*

Sergey Shekhovtsov, *Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

14.30 – 15.00

**PLENARY REPORT**

**Brain Asymmetries in the Cognitive Processing**

Boris M. Velichkovsky

*Corresponding Member of the Russian Academy of Science, Member of Academia Europaea*

15:00 – 15:30

**Keynote report**

**Non – stop trypanosomes**

Vyacheslav Yurchenko

*University of Ostrava, Ostrava, Czech Republic*

15:30 – 15:50

**Diversity and evolution of Tat LTR retrotransposon structures in non – flowering plants**

Mikhail Biryukov, Kirill Ustyantsev

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

15:50 – 16:10

**New data on Acanthobdellida phylogeny based on complete mitochondrial genomes**

Alexander Bolbat<sup>1</sup>, Gennadiy Vasiliev<sup>2</sup>, Irina Kaygorodova<sup>1</sup>, Vera Bogdanova<sup>2</sup>

<sup>1</sup>*Limnological Institute SB RAS, Irkutsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

16:10 – 16:30

**A genetic handicap approach: how to estimate the genome – wide burden of slightly – deleterious variants in a model population**

Victor Shamanskiy<sup>1</sup>, Konstantin Gunbin<sup>2</sup>, Konstantin Popadin<sup>1,3</sup>  
<sup>1</sup>*Immanuel Kant Baltic Federal University, Kaliningrad, Russia*  
<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
<sup>3</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*

**16:30 – 16:50**

**Sociality affects mutational spectrum of mtDNA in termites versus cockroaches**

Alina A. Mikhailova<sup>1,2</sup>, Thomas Bourguignon<sup>2</sup>, Konstantin Gunbin<sup>3</sup>, Konstantin Popadin<sup>1,4</sup>  
<sup>1</sup>*Immanuel Kant Baltic Federal University, Kaliningrad, Russia*  
<sup>2</sup>*Okinawa Institute of Science and Technology, Okinawa, Japan*  
<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
<sup>4</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*

**16:50 – 17:10**

**Coffee break**

**17:10 – 17:30**

**Mitochondrial mutational spectrum in poikilothermic versus homeothermic vertebrates: effects of the temperature**

Alina G. Mikhaylova<sup>1</sup>, Victor Shamanskiy<sup>1</sup>, Alina A. Mikhaylova<sup>1</sup>, Kristina Ushakova<sup>3</sup>, Konstantin Gunbin<sup>2</sup>, Konstantin Popadin<sup>1,4</sup>  
<sup>1</sup>*Immanuel Kant Baltic Federal University, Kaliningrad, Russia*  
<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*  
<sup>3</sup>*ITMO University, Saint Petersburg, Russia*  
<sup>4</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*

**17:30 – 17:50**

**The anatomy of mtDNA of mammals: the links with organismal longevity**

Victor Shamanskiy<sup>1</sup>, Kristina Ushakova<sup>1</sup>, Konstantin Popadin<sup>1,2</sup>, Konstantin Gunbin<sup>3</sup>  
<sup>1</sup>*Immanuel Kant Baltic Federal University, Kaliningrad, Russia*  
<sup>2</sup>*Ecole Polytechnique Federale de Lausanne, Lausanne, Switzerland*  
<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**17:50 – 18:10**

**CryProcessor: a novel tool for mining Cry toxins in Bacillus thuringiensis sequencing data**

Kirill Antonets<sup>1,2</sup>, Anton Shikov<sup>1,2</sup>, Yuri Malovichko<sup>1,2</sup>, Rostislav Skitchenko<sup>3</sup>, Anton Nizhnikov<sup>1,2</sup>  
<sup>1</sup>*All – Russia Research Institute for Agricultural Microbiology, St. Petersburg, Russia*  
<sup>2</sup>*Saint Petersburg State University, St. Petersburg, Russia*  
<sup>3</sup>*ITMO University, St. Petersburg, Russia*

**18:10 – 18:30**

**Mitochondrial genetics of amphipods: revealing mechanisms of diversity**

Elena V. Romanova<sup>1</sup>, Maria D. Logacheva<sup>2,3</sup>, Yuriy S. Bukin<sup>1,4</sup>, Elena A. Sirotinina, Dmitry Yu. Sherbakov<sup>1,4</sup>, Kirill V. Mikhailov<sup>2,3</sup>, Vladimir V. Aleoshin<sup>2,3</sup>  
<sup>1</sup>*LIN SB RAS, Irkutsk, Russia*  
<sup>2</sup>*Moscow State University, Moscow, Russia*

<sup>3</sup>IITP RAS, Moscow, Russia

<sup>4</sup>Irkutsk State University, Irkutsk, Russia

**18:30 – 18:50**

**Inter – vs. intraspecific genetic variability of morphologically similar ligophores species**

Ekaterina Vodiasova<sup>1</sup>, Alexei Ermolenko<sup>2</sup>, Evgenija Dmitrieva<sup>1</sup>, Dmitry Atopkin<sup>2</sup>, Olga Shikhat<sup>1</sup>

<sup>1</sup>IBSS RAS, Sevastopol, Russia

<sup>2</sup>FSC the East Asia Terrestrial Biodiversity, Vladivostok, Russia

**18:50 – 19:10**

**Distribution of Runs Of Homozygosity (ROHs) along the human genome is shaped by recombination and purifying selection**

K. Popadin<sup>1</sup>, E. Zezyulya<sup>2</sup>, A. Raymond<sup>3</sup>, D. Iliushchenko<sup>2</sup>

<sup>1</sup>EPFL, Lausanne, Switzerland

<sup>2</sup>IKBFU, Kaliningrad, Russia

<sup>3</sup>University of Lausanne, Lausanne, Switzerland

## Poster session

**Genetic diversity of the flat leeches (Hirudinea, Glossiphoniidae) in Western Siberia**

Nadezhda Bolbat<sup>1</sup>, Lyudmila Fedorova<sup>2</sup>, Irina Kaygorodova<sup>3</sup>

<sup>1</sup>Irkutsk State University, Irkutsk, Russia

<sup>2</sup>Surgut State University, Surgut, Russia

<sup>3</sup>Limnological institute SB RAS, Irkutsk, Russia

**Genetic aspects of internet – dependence in teenagers**

Marina Smolnikova, Sergey Tereshchenko

Scientific Research Institute of Medical Problems of the North FRC KSC SB RAS, Krasnoyarsk, Russia

**Application of ITS1 and ITS2 for population genetic studies of sturgeons (Acipenseridae)**

Guzel Davletshina<sup>1,2</sup>, Sergey Kliver<sup>1</sup>, Dmitry Prokopov<sup>1</sup>, Elena Interesova<sup>3</sup>, Vladimir Trifonov<sup>1,4</sup>

<sup>1</sup>IMCB SB RAS, Novosibirsk, Russia

<sup>2</sup>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

<sup>3</sup>TSU, Tomsk, Russia

<sup>4</sup>NSU, Novosibirsk, Russia

**Distribution of Bax protein in the rat hippocampus**

Pavel Lisachev, Anna Proskura

Institute of Computational Technologies, SB RAS, Novosibirsk, Russia

## **Methylation and expression profiles in Apoe vicinity point to specific neighboring interaction of Apoe and TOMM40 genes: implication for The Alzheimer disease**

Vladimir Babenko

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

## **EVALUATION OF *Sinorhizobium meliloti* GENOMIC ISLANDS INSERTED INTO THE tRNA – Thr**

Mariia Vladimirova, Alexey Afonin | Viktoria Muntyan | Boris Simarov | Marina Roumiantseva

*ARRIAM, Saint Petersburg, Russia*

## **Bioinformatic basis for species formation in the bacterial genera *Pectobacterium* and *Dickeya***

Peter Evseev<sup>1</sup>, Alexander Ignatov<sup>2,3</sup>, Konstantin Miroshnikov<sup>1</sup>

<sup>1</sup>*IBCh RAS, Moscow, Russia*

<sup>2</sup>*Research Center “PhytoEngineering” Ltd., Rogachevo, Moscow region,*

<sup>3</sup>*RUDN, Moscow, Russia*

## **Phylostratigraphic approach in evolutionary analysis: comparison of methods**

Tatiana Martusheva<sup>1</sup>, Zakhar Mustafin<sup>2</sup>, Sergey Lashin<sup>1,2</sup>

<sup>1</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>2</sup>*Kurchatov Genomics Center, ICG SB RAS, Novosibirsk, Russia*

## **Novel archaeal metagenome assembled genomes from acidophilic microbial community of Parys Mountain copper mine (UK)**

Aleksei Korzhenkov<sup>1</sup>, Stepan V. Toshchakov<sup>2</sup>, Ilya V. Kublanov<sup>2</sup>, Peter N. Golyshin<sup>3</sup>, | Olga V. Golyshina<sup>3</sup>

<sup>1</sup>*Kurchatov genome center, NRC Kurchatov Institute, Moscow, Russia*

<sup>2</sup>*Winogradsky Institute of Microbiology FRC “Biotechnology” RAS, Moscow, Russia*

<sup>3</sup>*Bangor University, Bangor, UK*

## **Dynamics and hypotheses of gene order shifts in mitochondrial genomes of Baikalian amphipods**

Elena A. Sirotinina<sup>1</sup>, Elena V. Romanova<sup>1</sup>, Dmitry Yu. Sherbakov<sup>1,2</sup>

<sup>1</sup>*LIN SB RAS, Irkutsk, Russia*

<sup>2</sup>*Irkutsk State University, Irkutsk, Russia*

## **Comparative Genomic Analysis of Moderate Bacteriophages of Alfalfa Root Nodule Bacteria**

Muntyan V.S.<sup>1</sup>, Muntyan A.N.<sup>1</sup>, Antonova E.V.<sup>2</sup>, Kozlova A.P.<sup>1</sup>, Dzyubenko E.A.<sup>3</sup>, Roumiantseva M.L.<sup>1</sup>, Afonin A.M.<sup>1</sup>, Kabilov M.R.<sup>4</sup>

<sup>1</sup>*All – Russian research institute for agricultural microbiology, Pushkin, Saint – Petersburg, Russia*

<sup>2</sup>*Institute of Plant and Animal Ecology, Ural Division of Russian Academy of Sciences,*

<sup>3</sup>*Federal Research Center N. I. Vavilov All – Russian Institute of Plant Genetic Resources (VIR) Ministry of science and higher education, Saint – Petersburg, Russia*

<sup>4</sup>*ICBFM SB RAS, Novosibirsk, Russia*

### **Plastid genome evolution in the genus *Allium***

Victoria Scobeyeva<sup>1,2</sup>, Denis Omelchenko<sup>3</sup>, Maria Logacheva<sup>1,4</sup>, Maxim Antipin<sup>1</sup>, Ilya Artyushin<sup>1</sup>, Andrey Samoilo<sup>5</sup>, Evgenii Konorov<sup>6</sup>, Maxim Belenikin<sup>2</sup>, Anastasiya Krinitsina<sup>1</sup>, Anna Speranskaya<sup>1,5</sup>

<sup>1</sup>*Lomonosov Moscow State University, Moscow, Russia*

<sup>2</sup>*Moscow Institute of Physics and Technology, Moscow region, Russia*

<sup>3</sup>*Institute for Information Transmission Problems, Moscow, Russia*

<sup>4</sup>*Skolkovo Institute of Science and Technology, Moscow, Russia*

<sup>5</sup>*Central Research Institute of Epidemiology, Moscow, Russia*

<sup>6</sup>*Vavilov Institute of General Genetics RAS, Moscow, Russia*

### **Intraspecific genetic variability of enhancers in the craniofacial tissue**

Elena Minkina<sup>1</sup>, Natalia Akberova<sup>1</sup>, Elena Shagimardanova<sup>1</sup>, Igor Adameyko<sup>2,3</sup>, Oleg Gusev<sup>1,4</sup>

<sup>1</sup>*Institute of Fundamental Medicine and Biology, KFU, Kazan, Russia*

*minkinaea@gmail.com*

<sup>2</sup>*Karolinska Institutet, Stockholm, Sweden*

<sup>3</sup>*Medical University Vienna, Vienna, Austria*

<sup>4</sup>*RIKEN, Yokohama, Japan*

### **Analysis of sequenced chromosome – specific libraries of gekkonids sheds light to large scale genome reshuffling in reptiles**

Katerina Tishakova<sup>1,2</sup>, Dmitry Prokopov<sup>1</sup>, Ilya Kichigin<sup>1</sup>, Anna Molodtseva<sup>1</sup>, Lukáš Kratochvíl<sup>3</sup>, Artem Lisachov<sup>4</sup>, Vladimir Trifonov<sup>1,2</sup>

<sup>1</sup>*Institute of Molecular and Cellular Biology SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*Charles University, Prague, Czech Republic*

<sup>4</sup>*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

### **Multigene phylogenies for the earthworm *Eisenia nordenskioldi* (Lumbricidae, Annelida)**

Sergei V. Shekhovtsov, Alexandra A. Shipova, Tatiana V. Polyboyarova, Sergei E. Peltek

*ICG SB RAS, Novosibirsk, Russia*

### **Genomic analyses of *Novymonas esmeraldas* and *Ca. Pandoraea novymonadis***

Alexandra Zakharova<sup>1</sup>, Daria Tashyreva<sup>2</sup>, Jorge Morales<sup>3</sup>, Eva Nowack<sup>3</sup>, Julius Lukeš<sup>2</sup>, Vyacheslav Yurchenko<sup>1</sup>

<sup>1</sup>*University of Ostrava, Ostrava, Czech Republic*

<sup>2</sup>*Institute of Parasitology Biological Centre, České Budějovice, Czech Republic*

<sup>3</sup>*Heinrich Heine University, Düsseldorf, Germany*

### **Genome and Karyotype Evolution after Whole Genome Duplication in Free – Living Flatworms of the Genus *Macrostomum***

Kira Zadesenets<sup>1</sup>, Nikita Ershov<sup>1</sup>, Dmitry Oshchepkov<sup>1</sup>, Eugene Berezikov<sup>1,2</sup>, Lukas Schärer<sup>3</sup>, Nikolay B. Rubtsov<sup>1</sup>

<sup>1</sup>*ICG SB RAS, Novosibirsk, Russia*

<sup>2</sup>*ERIBA, Groningen, The Netherlands*

<sup>3</sup>University of Basel, Basel, Switzerland

### **Comparative genomics of heat shock proteins system in extremophile nonbiting midges**

Olga Kozlova | Guzel Gazizova | Elena Shagimardanova | Oleg Gusev  
*Kazan Federal University, Kazan, Russia*

### **The phenotypic manifestation of Wolbachia genetic diversity in host fitness**

Elena V. Burdina, Nataly Gruntenko, Petr Menshanov, Roman Bykov, | Yury Ilinsky, Inga Rauschenbach  
*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

### **FMO superfamily protein phylogeny and the origin of YUCCA family.**

Igor Turnaev, Valentin Suslov, Konstantin Gunbin, Dmitriy Afonnikov  
*Institute of Cytology and Genetics, ICG SB RAS, Novosibirsk, Russia*

### **Phylogenetic Analysis of Poxviridae Genomes Using K – mer Approach**

Tatyana Nepomnyashchikh, Denis Antonets, Tatyana Tregubchak, Alexander Shvalov, Elena Gavrilova, Rinat Maksyutov  
*SRC VB "Vector" Rospotrebnadzor, Koltsovo, Russia*

### **Candidate SNP markers of rheumatoid arthritis changing the affinity of TATA – binding protein for the human gene promoters expo disruptive selection of immunoactivative and immunosuppressive genes that provoke and prevent this disorder, respectively, as if it could be a self – domestication syndrome**

Dmitry Rasskazov<sup>1</sup>, Irina Chadaeva<sup>1</sup>, Mikhail Ponomarenko<sup>1</sup>, Ekaterina Sharypova<sup>1</sup>, Irina Drachkova<sup>1</sup>, Maria Nazarenko<sup>2</sup>

<sup>1</sup>*Institute of Cytology and Genetics, ICG SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Institute of Medical Genetics, IMG TNRM RAS, Tomsk, Russia*

### **Circular RNA host gene and orthologue prediction using the self – designed CircParser pipeline**

Artem Nedoluzhko<sup>1</sup>, Fedor Sharko<sup>2</sup>, Golam Rbanni<sup>1</sup>, Anton Teslyuk<sup>2</sup>, Ioannis Konstantinidis<sup>1</sup>, Jorge M.O. Fernandes<sup>1</sup>

<sup>1</sup>*Nord University, Bodø, Norway*

<sup>2</sup>*NRC "Kurchatov Institute", Moscow, Russia*

### **Hydroxymethylation changes during early embryonic development in zebrafish**

Artem Nedoluzhko<sup>1</sup>, Paula Berrutti<sup>1</sup>, Igo Guimarães<sup>2</sup>, Ioannis Konstantinidis<sup>1</sup>, Igor Babiak<sup>1</sup>, Jorge M.O. Fernandes<sup>1</sup>

<sup>1</sup>*Nord University, Bodø, Norway*

<sup>2</sup>*Universidade Federal de Goiás, Goiás, Brazil*

### **OrthoWeb – web application for macro – and microevolutionary analysis of genes**

Zakhar Mustafin<sup>1</sup>, Alexey Mukhin<sup>1</sup>, Dmitry Afonnikov<sup>1,2</sup>, Yury Matushkin<sup>3</sup>, Sergey Lashin<sup>1,2</sup>

<sup>1</sup>*Kurchatov Genomics Center, ICG SB RAS, Novosibirsk, Russia*

<sup>2</sup>*NSU, Novosibirsk, Russia*

<sup>3</sup>*ICG SB RAS, Novosibirsk, Russia*

### **Genomic Signals of Adaptation in the Northern Ural and Western Siberian Populations**

Gennady Khvorykh<sup>1</sup>, Giang Vu<sup>2</sup>, Andrey Khrunin<sup>1</sup>

<sup>1</sup>*Institute of Molecular Genetics of the Russian Academy of Sciences, Moscow, Russia*

<sup>2</sup>*Moscow Polytechnic University, Moscow, Russia*

# SYMPOSIUM

## Animal Genetics and genomics

**Oral reports**

10 July, Friday

**Small Conference Hall**

**Morning session. Animal Genetics and genomics**

**Chairs: Mikhail Moshkin, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia**

**Nikolai Yudin, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia**

**Vladimir Naumenko, Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia**

9.00 – 9.30

**PLENARY REPORT**

**Mathematical problems driven by COVID-19**

*Sergey Kabanikhin*

*Novosibirsk State University, Novosibirsk, Russia*

9:30 – 10:10

**Keynote report**

**Update status of mouse resources for studies of gene function and disease at RIKEN BRC**

*Atsushi Yoshiki*

*RIKEN BioResource Research Center and Head of Experimental Animal Division, Tsukuba, Ibaraki, Japan*

10:10 – 10:40

**Posters show**

10:40 – 11:05

**Differentially Expressed Genes in Longitudinal Axis of the Fox's Hippocampus**

*Yury Alexandrovich, Larisa Meister, Yury Herbeck*

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

11:05 – 11:30

**Genetic structure of breeding pigs of Large White, bred in Russia**

*Lyubov Getmantseva, S. Bakoev, O. Kostyunina, A. Traspov, Yu. Prytkov, N. Bakoev*

*Federal Science Center for Animal Husbandry named after Academy Member L.K. Ernst Dubrovica, Russia*

11:30 – 11:50

**Rapidly evolving SNPs feature highly significant trait associations in GWAS SNP hotspots**

*Roman Babenko, Anton Zhuravlev*

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**Lunch**

**Evening session. Animal Genetics and genomics**

**Chairs: Mikhail Moshkin, *Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia***

**Nikolai Yudin, *Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia***

**Vladimir Naumenko, *Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia***

14:30 – 15:00

**PLENARY REPORT**

**Epidemics: challenges and responses**

*Alexey Romanykha*

*Marchuk Institute of Numerical Mathematics, Moscow, Russia*

<b>15:00 – 15:30</b>	<b>Population genetic variation of serotonin transporter gene (SLC6A4), associated with neurophysiological development</b> Shyamala Hande <i>Melaka Manipal Medical College, Manipal, India</i>
<b>15:30 – 15:50</b>	<b>Hippocampal Overexpression of The Cerebral Dopamine Neurotrophic Factor (CDNF) Impaired Fear Memory Formation in Rats</b> Tatiana Ilchibaeva, Elizaveta Zolotenkova, Dmitry Eremin, <u>Anton Tsybko</u> <i>Institute of Cytology and Genetics, SB RAS, Nobosibirsk, Russia</i>
<b>15:50 – 16:10</b>	<b>The First Evidences of Direct Interaction Between 5 – HT2A and TrkB receptors</b> <u>Tatiana Ilchibaeva</u> , Anton Tsybko, Vladimir Naumenko <i>Institute of Cytology and Genetics SB RAS, Nobosibirsk, Russia</i>
<b>16:10 – 16:40</b>	<b>Testing inter – relations between disturbed sleep and sterility in intra – specific hybrids of fruit fly</b> Lyudmila Zakharenko <sup>1</sup> , Dmitriy Petrovskii <sup>1</sup> , <u>Arcady Putilov</u> <sup>2</sup> <sup>1</sup> <i>Institute of Cytology and Genetics, SB RAS, Nobosibirsk, Russia</i> <sup>2</sup> <i>Federal Research Centre for Fundamental and Translational Medicine, Novosibirsk, Russia</i>
<b>16:40 – 17:10</b>	<b>Coffee break</b>
<b>17:20 – 17:50</b>	<b>Effects of long – term ethanol consumption in mice: interaction between BDNF and brain serotonin systems</b> <u>Vladimir Naumenko</u> , Tatiana Ilchibaeva, Egor Antonov, Darya Bazovkina, Nina Popova <i>Institute of Cytology and Genetics, SB RAS, Nobosibirsk, Russia</i>
<b>17:50 – 18:10</b>	<b>Histological evaluation of postnatal retinal development of senescence – accelerated OXYS rats</b> <u>Darya Telegina</u> <sup>1</sup> , Anna Antonenko <sup>2</sup> , Oyuna Kozhevnikova <sup>1</sup> <sup>1</sup> <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i> <sup>2</sup> <i>Novosibirsk State University, Novosibirsk, Russia</i>
<b>18:10 – 18:30</b>	<b>Mechanisms and functions of neurogenesis in the limbic system of adult animals</b> Tatyana Zapara <sup>1</sup> , Alexander Romashchenko <sup>2</sup> , Anna Proskura <sup>1</sup> , Alexander Ratushnyak <sup>1</sup> , Svetlana Vechkapova <sup>1</sup> <sup>1</sup> <i>Institute of Computational Technologies, SB RAS, Novosibirsk, Russia</i> <sup>2</sup> <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i>
<b>18:30 – 18:50</b>	<b>Effect of overexpression of the 5 – HT7 receptor gene on behavior and brain serotonin system in ASC mice with predisposition to depressive – like behavior</b> Irina Baraboshkina, Darya Bazovkina, Tatiana Ilchibaeva, Egor Antonov, Elizabeth Kulikova, Vladimir Naumenko <i>Institute of Cytology and Genetics, SB RAS, Nobosibirsk, Russia</i>

**Diversity and evolution of Tat LTR retrotransposon structures in non – flowering plants**

Mikhail Biryukov, Kirill Ustyantsev

Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia

**A platform for storage and analysis of results of genome – wide association studies of sheep**

Alexander S. Zlobin<sup>1</sup>, Anatoliy V. Kirichenko<sup>1</sup>, Tatyana I. Shashkova<sup>2</sup>, Natalya A. Volokova<sup>3</sup>, Pavel M. Borodin<sup>3</sup>, Lennart C. Karsen<sup>4</sup>, Yakov A. Tsepilov<sup>1</sup>, Yurii S. Aulchenko<sup>1</sup>

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*L.K. Ernst Federal Science Center for Animal Husbandry, Dubrovitsy, Moscow Region, Russia*

<sup>4</sup>*PolyOmica, 's – Hertogenbosch, the Netherlands*

**Effect of bispidine containing monoterpene moieties on physical performance in mice**

Anastasiya A. Kotlyarova<sup>1</sup>, Dina V. Korchagina<sup>2</sup>, Tatyana G. Tolstikova<sup>2</sup>, Konstantin Yu. Ponomarev<sup>2</sup>

<sup>1</sup>*Research Institute of Clinical and Experimental Lymphology – a branch of the Institute of Cytology and Genetics of Siberian Branch of Russian Academy of Sciences;* <sup>2</sup>*Novosibirsk Institute of Organic Chemistry of Siberian Branch of Russian Academy of Sciences, Novosibirsk, RUSSIA*

**Influence of the factors of maternal milieu on taste preferences and metabolic parameters in mouse male and female offspring**

Elena Denisova<sup>1</sup>, Elena Makarova<sup>1</sup>, Maria Savinkova<sup>2</sup>

<sup>1</sup>*Institute of Cytology and Genetics SB RAS;* <sup>2</sup>*Novosibirsk State University, Novosibirsk, RUSSIA*

**Genetic aspects of internet – dependence in teenagers**

Marina Smolnikova, Sergey Tereshchenko

*Scientific Research Institute of Medical Problems of the North FRC KSC SB RAS, Krasnoyarsk, Russia*

**The cross – talk molecular pathways of glutamate and leptin receptors**

Anna L. Proskura, Mariya Yu. Islamova, Svetlana O. Vechkapova

*Institute of Computational Technologies, SB RAS, Novosibirsk, Russia*

**Accumulation of oxidative hepatobiliary lesions during experimental opisthorchiasis**

Oxana Zaparina<sup>1</sup>, Anna Kovner<sup>1</sup>, Maria Pakharukova<sup>1,2</sup>, Viacheslav Mordvinov<sup>1</sup>

<sup>1</sup>*ICG SB RAS, Novosibirsk, RF;*

<sup>2</sup>*NSU, Novosibirsk, RF*

**Enlarged clinical Belarusians' exomes: opportunities and restrictions of additional analysis**

Aleh Liaudanski, Danat Yermakovich

*IGC NAS, Minsk, Belarus*

**Multivariate analysis identify new loci associated with meat productivity and carcass traits in sheeps (Ovis aries)**

Alexander S. Zlobin<sup>1</sup>, Natalia A. Volkova<sup>2</sup>, Pavel M. Borodin<sup>2</sup>, Tatiana I. Aksenovich<sup>2</sup>, | Yakov A. Tsepilov<sup>3</sup>

<sup>1</sup>Kurchatov Genomic Center of IC&G, Novosibirsk, Russia

<sup>2</sup>L.K. Ernst Federal Science Center for Animal Husbandry, Dubrovitsy, Moscow Region, Russia

<sup>3</sup>Novosibirsk State University, Novosibirsk, Russia

**Study of the COI Gene Fitness for a Population – Genetic Analysis of Endemic Baikal Sponges *L. Baikalensis***

Alena Yakhnenko<sup>1,2</sup>, Valeria Itskovich<sup>1</sup>

<sup>1</sup>LIN SB RAS, Irkutsk, Russia

<sup>2</sup>JINR, Dubna, Russia

**Metabolic response of the Siberian frog *Rana amurensis* to anoxia**

Sergei V. Shekhovtsov<sup>1</sup>, Nina A. Bulakhova<sup>2</sup>, Yuri P. Tsentalovich<sup>3</sup>, Ekaterina A. Zelentsova<sup>3</sup>, Daniil I. Berman<sup>2</sup>

<sup>1</sup>ICG SB RAS, Novosibirsk, Russia; <sup>2</sup>IBPN FEB RAS, Magadan, Russia; <sup>3</sup>ITC SB RAS, Novosibirsk, RUSSIA

**MtDNA variability in the field vole (*Microtus agrestis* L., 1761), Arvicolinae, Rodentia) in the Urals and adjacent territories**

Maria Krokhalova, Lidia Yalkovskaya, Petr Sibiryakov, Evgenia Markova, Aleksandr Borodin

*Institute of Plant and Animal Ecology, UB RAS, Ekaterinburg, Russia*

# Symposium "Systems biology of DNA repair processes and programmed cell death"

Oral reports

7 July, Tuesday

Computer Class

Evening session 1. Systems biology of DNA repair processes and programmed cell death

Chair: Inna Lavrik, *Otto von Guericke University, Magdeburg, Germany*

15:00 – 15:40

**Keynote report**

**Ferroptosis, a metabolic death pathway**

Marcus Conrad

*Helmholtz Zentrum München, Neuherberg, Germany.*

15:40 – 16:20

**Keynote report**

**TRAIL-resistance in melanoma: What doesn't kill you makes you move!**

Dagmar Kulms

*Technische Universität Dresden, Dresden, Germany*

16:20 – 16:45

**Targeting CD95 signaling network**

Inna Lavrik,

<sup>1</sup>*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Translational Inflammation Research, Medical Faculty, Otto von Guericke University*

*Magdeburg, Magdeburg, Germany*

16:45 – 17:00

**Coffee break**

17:00 – 17:25

**Computational insights into molecular mechanisms of CD95 programmed cell death activation**

Nikita Ivanisenko<sup>1</sup>, Vladimir A. Ivanisenko<sup>1</sup>, Laura K. Hillert<sup>2</sup>, Corinna König<sup>2</sup>, Inna N.

Lavrik<sup>1,2</sup>

<sup>1</sup>*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Translational Inflammation Research, Medical Faculty, Otto von Guericke University Magdeburg, Magdeburg, Germany*

17:25 – 17:45

**Mitophagy promotes cell death pathways induced by lactaptin**

Fabian Wohlfromm

*Translational Inflammation Research, Medical Faculty,*

*Velichkovskii*

*, Magdeburg, Germany*

17:45– 18:30

**Posters show**

**Oral reports**

10 July, Friday

**Library**

**Evening session 2. Systems biology of DNA repair processes and programmed cell death**

**Chair: Olga Lavrik, *Institute of Chemical Biology and Fundamental Medicine of SB RAS, Novosibirsk, Russia*  
Dmitry Zharkov, *Institute of Chemical Biology and Fundamental Medicine of SB RAS, Novosibirsk, Russia***

14:30 – 15:00

**PLENARY REPORT**

**Epidemics: challenges and responses**

Alexey Romanykha

*Marchuk Institute of Numerical Mathematics, Moscow, Russia*

<b>15:00 – 15:30</b>	<p><b>PARP1 activation directs RNA binding proteins to DNA damages to form PARG reversible compartments enriched in damaged DNA</b></p> <p>Olga Lavrik<sup>1</sup>, Maria Sukhanova<sup>1</sup>, Anastasia Singatulina<sup>1</sup>, Konstantin Naumenko<sup>1</sup>, Loic Hamon<sup>2</sup>, David Pastré<sup>2</sup></p> <p><sup>1</sup><i>Institute of Chemical Biology and Fundamental Medicine SB RAS, Novosibirsk, Russia</i></p> <p><sup>2</sup><i>Université Paris – Saclay, Evry, France</i></p>
<b>15:30 – 16:00</b>	<p><b>Human apurinic/apyrimidinic endonuclease 1 is modified by poly(ADP – ribose) polymerase 1 via the DNA structure – controlled mechanism</b></p> <p>Nina Moor, Inna Vasil'eva, Nikita Kuznetsov, Olga Lavrik</p> <p><i>Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia</i></p>
<b>16:00 – 16:25</b>	<p><b>Nucleotide Excision Repair Proteins and PARP1/PAR Interplay Regulates Protein Assembly on Damaged DNA</b></p> <p>Nadejda Rechkunova, Maria Sukhanova, Ekaterina Maltseva, Olga Lavrik, Yuliya Krasikova</p> <p><i>Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia</i></p>
<b>16:25 – 16:50</b>	<p><b>Helicase XPD <i>Chaetomium thermophilum</i> as a functional analogue of human XPD</b></p> <p>Irina Petrusheva<sup>1</sup>, Natalia Lukyanchikova<sup>1</sup>, Olga Lavrik<sup>1</sup>, Jochen Kuper<sup>2</sup>, Janette Kappenberger<sup>2</sup>, Rashid Anarbaev<sup>1</sup>, Caroline Kisker<sup>2</sup></p> <p><sup>1</sup><i>Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia</i></p> <p><sup>2</sup><i>Rudolf Virchow Center for Experimental Medicine, University Wurzburg, Germany</i></p>
<b>16:50 – 17:10</b>	<b>Coffee break</b>
<b>17:10 – 17:50</b>	<p><b>Actors of the base excision repair play: How well do we know the credits?</b></p> <p>Dmitry O. Zharkov<sup>1,2</sup>, Anton V. Endutkin<sup>2</sup>, Evgeniia A. Diatlova<sup>2</sup>, Anna V. Yudkina<sup>1,2</sup>, Alexander V. Popov<sup>2</sup></p> <p><sup>1</sup><i>Novosibirsk State University, Novosibirsk, Russia</i></p> <p><sup>2</sup><i>Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia</i></p>
<b>17:50 – 18:20</b>	<p><b>Initial steps of base excision repair on DNA – substrates with non – canonical structures</b></p> <p>Alexandra A. Kuznetsova, Olga S. Fedorova, Anastasiia T. Davletgildeeva, Nikita A. Kuznetsov</p> <p><i>Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia</i></p>
<b>18:20 – 18:45</b>	<p><b>The role of DNA repair in active DNA demethylation is studied by the construct based on the CRISPR/Cas9 system</b></p> <p>Zarina Kakhkharova, Darya Khantakova, Inga Grin</p> <p><i>Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia</i></p>
<b>18:45 – 19:10</b>	<p><b>Conformational dynamics in methylated DNA repair by human Fe(II)/alpha – ketoglutarate dependent dioxygenases ALKBH2 and ALKBH3</b></p> <p>Lyubov Yu. Kanazhevskaya<sup>1</sup>, Denis A. Smyshlyaev<sup>1,2</sup>, Olga S. Fedorova<sup>1</sup></p> <p><sup>1</sup><i>Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia</i> <sup>2</sup><i>Novosibirsk State University</i></p>

**Activity of human AP – endonuclease APE1 on DNA – and RNA – substrates forming non – canonical structures**

Anastasiia T. Davletgildeeva, Olga S. Fedorova, Alexandra A. Kuznetsova, Nikita A. Kuznetsov  
*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

**The effect of protein – protein interactions on the activity of APE1 SNP forms**

Olga A. Kladova, Nikita A. Kuznetsov, Irina V. Alekseeva, Olga S. Fedorova  
*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

**Activity of SNP variants of human uracil – DNA glycosylases SMUG1 and MBD4**

Irina V. Alekseeva, Nikita A. Kuznetsov, Artemiy S. Bakman, Olga S. Fedorova  
*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

**Activity of DNA glycosylases on non – canonical DNA substrates**

Evgeniia Diatlova, Dmitry Zharkov  
*Novosibirsk State University, Novosibirsk, Russia*  
*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

**Inhibition of DNA – repairing enzymes by nucleoside derivatives**

Mikhail S. Drenichev<sup>1</sup>, Alexandra L. Zakharenko<sup>2</sup>, Nadezhda S. Dyrkheeva<sup>2</sup>, Georgy A. Ivanov<sup>1</sup>, Vladimir E. Oslovsky<sup>1</sup>, Ekaterina S. Ilina<sup>2</sup>, Irina A. Chernyshova<sup>2</sup>, Olga I. Lavrik<sup>2</sup>, Sergey N. Mikhailov<sup>1</sup>  
<sup>1</sup> *Engelhardt Institute of Molecular Biology, Russian Academy of Sciences, Moscow, Russia*  
<sup>2</sup> *Institute of Chemical Biology and Fundamental Medicine SB RAS, Novosibirsk, Russia*

**Lesion recognition and cleavage of damage – containing G – quadruplexes by DNA glycosylases**

Aleksandra A. Kuznetsova, Olga S. Fedorova, Nikita A. Kuznetsov  
*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

**Nucleosome assembling: quick – time reconstitution protocol**

Alexander Ukraintsev, Ekaterina Belousova, Michael Kutuzov, Svetlana Khodyreva, Tatyana Kurgina, Olga Lavrik  
*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

**PARP1 activation promotes FUS translocation to cytoplasm and incorporation into stress granules**

Anastasia Shavkatovna Singatulina<sup>1</sup>, Bénédicte Desforges<sup>2</sup>, Pastré David<sup>2</sup>, Maria Vladislavovna Sukhanova<sup>1</sup>, Ahmed Bouhss<sup>2</sup>, Loic Hamon<sup>2</sup>, Olga Ivanovana Lavrik<sup>1</sup>  
<sup>1</sup> *Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*  
<sup>2</sup> *Université Paris – Saclay, Evry, France*

**Platinum Polyoxoniobates have potential as anticancer agents**

Anna V. Yudkina<sup>1,2</sup>, Pavel A. Abramov<sup>3</sup>, Ivan P. Vokhtantsev<sup>1,2</sup>, Inga R. Grin<sup>1,2</sup>, Maxim N. Sokolov<sup>3</sup>, Dmitry O. Zharkov<sup>1,2</sup>  
<sup>1</sup> *Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*Nikolaev Institute of Inorganic Chemistry SB RAS, Novosibirsk, Russia*

### **The influence of ligand structure of ruthenium nitrosyl complexes on their biological activity**

Darya Khantakova<sup>1,2</sup>, Inga Grin<sup>1,2</sup>

<sup>1</sup>*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

### **Single – nucleotide polymorphisms of hNEIL2 gene: from protein structure to functions in base excision DNA repair**

Zarina Kakhkharova<sup>1,2</sup>, Petrova Daria<sup>1,2</sup>, Inga Grin<sup>1,2</sup>

<sup>1</sup>*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

### **The interplay between NHEJ and BER in NHEJ deficient cells**

Polina Loshchenova<sup>1,2</sup>, Svetlana Sergeeva<sup>1,2</sup>, Grigory Dianov<sup>1,2,3</sup>

<sup>1</sup>*Institute of Chemical Biology and Fundamental Medicine SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University*

<sup>3</sup>*Oxford Institute for Radiation Oncology, University of Oxford, UK*

### **Processing of Clustered DNA Damages by Nucleotide Excision Repair pathway**

Natalia Lukianchikova, Petruseva Irina, Alexander Lomzov, Olga Lavrik

<sup>1</sup>*CBFM SB RAS, Novosibirsk, Russia*

### **YB – 1 as modulator of PARP1 activity**

K.N. Naumenko, M.V. Sukhanova, E.E. Alemasova, T.A. Kurgina, M.M. Kutuzov, O.I. Lavrik

*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

### **Sensitization mechanism of cells with TDP1 inhibitors to the action of topotecan**

Nadezhda S. Dyrkheeva<sup>1</sup>, Irina V. Il'ina<sup>2</sup>, Nikolay S. Li – Zhulanov<sup>2</sup>, Anastasiya A. Malakhova<sup>3</sup>, Sergey P. Medvedev<sup>3</sup>, Suren M. Zakian<sup>3</sup>, Konstantin P. Volcho<sup>2</sup>, Nariman F. Salakhutdinov<sup>2</sup>, Olga I. Lavrik<sup>1</sup>

<sup>1</sup>*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*N.N. Vorozhtsov Novosibirsk Institute of Organic Chemistry, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

### **Computer-assisted analysis of caspases molecular evolution**

Alexey Zamaraev<sup>1</sup>, Gelina Kopeina<sup>1</sup>, Boris Zhivotovsky<sup>1,2</sup>, Konstantin Gunbin<sup>3</sup>

<sup>1</sup>*MSU, Moscow, Russia*

<sup>2</sup>*Karolinska Institutet, Stockholm*

<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**PARP1 and PARP2 affinity to the lesions in the context of nucleosomes**

Tatiana Andreevna Kurgina, Rashid Oktamovich Anarbaev, Ekaterina Anatolyevna Belousova, Olga Ivanovana Lavrik, Michail Michailovich Kutusov, Svetlana Nikolaevna Khodireva  
*Institute of Chemical Biology and Fundamental Medicine, SB RAS, Novosibirsk, Russia*

# SYMPOSIUM

## Cognitive Science and Genomics

Oral reports

9 July, Thursday

Library

Morning session. Cognitive Science and Genomics

Chair: Tamara Amstislavskaya, *PhBMRI, Novosibirsk, Russia*

9.00 – 9.30

**PLENARY REPORT**

**Genetic Development of the Nociceptive System and the Epigenetics of the Experience of Pain**

Jack M. Berger

*Keck School of Medicine of USC, University of Southern California, Los Angeles, California, USA*

9:30 – 10:00

**Keynote report**

**ENIGMA: The Quest for Genetic Loci that Affect the Speed of Brain Development and Aging in 50,000 People from 45 Countries**

Paul M. Thompson, *for the ENIGMA Consortium,*

*University of Southern California, Los Angeles, CA, USA*

10:00 – 10:30

**Keynote report**

**Pharmacological effects of arecoline on zebrafish behavior, neurochemistry, neurophysiology and brain gene expression**

Tamara Amstislavskaya<sup>1</sup>, Nazar Serikul<sup>2</sup>, Erik Alpyshov<sup>2</sup>, DongMei Wang<sup>2</sup>, JingTao Wang<sup>2</sup>, Allan Kalueff<sup>2,3</sup>

<sup>1</sup>*Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

<sup>2</sup>*School of Phar Southwest University, Chongqing, China*

<sup>3</sup>*Institute of Translational Biomedicine, St. Petersburg State University, St. Petersburg, Russia*

10:30 – 10:50

**An approach to the analysis of cognitive systems through the evolution of simple**

Aleksander Ratushnyak, Iliya Malakhin, Tatyana Zapara

*Institute of Computational Technologies, SB RAS, Novosibirsk, Russia*

10:50 – 11:05

**Diabetes Type 2 as a Risk Factor of Neurodegeneration Development and Cognitive Impairment in db/db Mice**

Tatiana Korolenko<sup>1</sup>, Nina Dubrovina<sup>1</sup>, Marina Ovsyukova<sup>1</sup>, Natalya Bgatova<sup>2</sup>, Alexander Pupyshev<sup>1</sup>, Elena Anufrienko<sup>1</sup>, Chih – Li Lin<sup>3</sup>, Evgeniy Zavjalov<sup>2</sup>

<sup>1</sup>*Scientific Research Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Institute of Medicine, Chung Shan Medical University, Taichung, Taiwan*

11:05 – 11:20

**Expression of autophagy genes and markers of inflammation in the brain in a transgenic mouse model of Parkinson's disease**

Victor M. Belichenko<sup>1</sup>, Anna A. Akopyan<sup>1</sup>, Maria A. Tikhonova<sup>1</sup>, Alexandra B. Shintyapina<sup>2</sup>, Tatiana A. Korolenko<sup>1</sup>, Larisa A. Fedoseeva<sup>3</sup>, Tamara G. Amstislavskaya<sup>1</sup>

<sup>1</sup>*Scientific Research Institute of Physiology and Basic Medicine  
Novosibirsk, Russia*

<sup>2</sup>*Federal Research Center for Basic and Translational Medicine  
Novosibirsk, Russia*

<sup>3</sup>*Federal Research Center "Institute of Cytology and Genetics"  
Novosibirsk, Russia*

11:20 – 11:40

**Positive effect of joint activation of mTOR – dependent and mTOR – independent pathways of autophagy in the treatment of two experimental models of neurodegeneration**

Alexander Pupyshev, Nina Dubrovina, Maria Tikhonova, Anna Akopyan, Marina Ovsyukova, Mikhail Tenditnik, Tatiana Korolenko  
*Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

11:40 – 11:50

**Coffee break**

11:50 – 12:10

**Effects of diets rich in plant polyphenols in mouse models of neurodegenerative disorders**

Maria Tikhonova<sup>1</sup>, Tamara Amstislavskaya<sup>1</sup>, Anna Akopyan<sup>1</sup>, Marina Ovsyukova<sup>1</sup>, Michael Tenditnik<sup>1</sup>, Elena Khlestkina<sup>2,3</sup>

<sup>1</sup>*Scientific Research Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

<sup>2</sup>*N.I. Vavilov All – Russian Research Institute of Plant Genetic Resources, St. Petersburg, Russia*

<sup>3</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**12:10 – 12:30**      **Dynamic regulation of murine cortical transcriptome by early – life stress: Impairment of myelination and cognitive functions**

Natalya Bondar, Anastasia Shulyupova, Polina Kisaretova, Nikita Ershov, Elena Antontseva, Tatiana Merkulova

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**12:30 – 12:50**      **Monoamine signaling gene networks unraveled in mouse social stress model**

Vladimir Babenko, Natalia Kudryavtseva

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**12:50 – 13:10**      **NRG1, PIP4K2A, and HTR2C contain possible genetic biomarkers of several clinical subphenotypes of depression and bipolar disorder**

Anastasia Levchenko

*Theodosius Dobzhansky Center for Genome Bioinformatics, Saint Petersburg State University, Saint Petersburg, Russia*

**Lunch**

### Evening session. Cognitive Science and Genomics

**Chair: Alexander Savostyanov, *Institute of Physiology and Basic Medicine, Novosibirsk, Russia;*  
*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia***

**14.30 – 15.00**      **PLENARY REPORT**

**Brain Asymmetries in the Cognitive Processing**

Boris M. Velichkovsky

*Corresponding Member of the Russian Academy of Science, Member of Academia Europaea*

**15:00 – 15:20**      **Electroencephalographic correlates of an insight**

Gennady Knyazev<sup>1</sup>, Andrey Bocharov<sup>1,2</sup>, Alexander Savostyanov<sup>1,2</sup>

<sup>1</sup>*Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk National Research State University, Novosibirsk, Russia*

**15:20 – 15:40**      **Are younger people sleepier than older people after missing bedtime and night sleep? It depends...**

Arcady Putilov, Olga Donskaya

*Research Institute for Molecular Biology and Biophysics of the Federal Research Centre for Fundamental and Translational Medicine, Novosibirsk, Russia*

<b>15:40 – 16:00</b>	<b>EEG correlates of strategies of emotional regulation during perception of emotional information</b> <u>Andrey Bocharov</u> <i>Institute of Physiology and Basic Medicine, Novosibirsk, Russia</i>
<b>16:00 – 16:20</b>	<b>Electroencephalographic reactions under conditions of recognition of emotional written language in people residing in different regions of Siberia</b> Alexander Savostyanov <sup>1,2</sup> , Sergey Tamozhnikov <sup>2</sup> , <u>Natalya Milakhina</u> <sup>1</sup> , Darya Bazovkina <sup>1</sup> , Alexandra Karpova <sup>3</sup> , Natalia Borisova <sup>3</sup> , Elena Afanaseva <sup>3</sup> <sup>1</sup> <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i> <sup>2</sup> <i>Institute of Physiology and Basic Medicine, Novosibirsk, Russia</i> <sup>3</sup> <i>North – Eastern Federal University, Yakutsk, Russia</i>
<b>16:20 – 16:40</b>	<b>Behavioral and EEG effects of meditation on executive control functions and speech recognition</b> Alexander Savostyanov <i>Institute of Physiology and Basic Medicine, Novosibirsk, Russia</i> <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i>
<b>16:40 – 17:00</b>	<b>Study of personal qualities and EEG activity in a stop signal paradigm in residents of northern regions</b> <u>Tatiana Astakhova</u> <sup>1</sup> , Alexander Saprigyn <sup>2</sup> , Sergey Tamozhnikov <sup>2</sup> , Alexandra Karpova <sup>3</sup> , Natalia Borisova <sup>3</sup> , Elena Afanaseva <sup>3</sup> , Alexander Savostyanov <sup>1,2,4</sup> <sup>1</sup> <i>Novosibirsk State University, Novosibirsk, Russia</i> <sup>2</sup> <i>Institute of Physiology and Basic Medicine, Novosibirsk, Russia</i> <sup>3</sup> <i>North – Eastern Federal University, Yakutsk, Russia</i> <sup>4</sup> <i>Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia</i>
<b>17:00 – 17:20</b>	<b>Poster Show</b>

### Poster session

#### Genetic aspects of internet – dependence in teenagers

Marina Smolnikova

*Research Institute for Medical Problems in the North, Krasnoyarsk, Russia*

#### Cognitive functions and polymorphism of the BDNF gene in patients with schizophrenia and healthy individuals

Anastasiia Boiko, Ekaterina Mikhailitskaya, Elena Kornetova, Svetlana Ivanova

*Mental Health Research Institute Tomsk NRMС, Tomsk, Russia*

#### The impact of early – life stress on the expression of genes associated with the formation of the myelin sheath of neurons in the prefrontal cortex of 15 – day – old male mice.

Anastasia Shulyupova, Arina Smelova, Vasiliy Reshetnikov, Natalya Bondar

*Institute of Cytology and Genetics SB RAS, Novosibirsk, Russia*

**Compulsive – like behaviors in DISC1 – mice**

*Nadezhda Chizhova<sup>1</sup>, Kristina Smirnova<sup>2,3</sup>*

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Novosibirsk State University, Novosibirsk, Russia*

<sup>3</sup>*Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

**Altered expression of genes Npas4 and Nr1d1 in adult female mice with history of early – life stress**

*Yuliya Ryabushkina, Vasily Reshetnikov, Natalya Bondar*

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

**Associations of polymorphic variants of the genes of neurotrophic factors BDNF, NGF, NRG1 with remission in patients with depressive disorders**

*Natalia Vyalova, German Simutkin, Nikolay Bokhan, Svetlana Ivanova*

*Mental Health Research Institute of TNRMC, Tomsk, Russia*

**Immunoglobulins with proteolytic activity as a biomarker of impaired humoral immune system in schizophrenia**

*Evgeny Ermakov<sup>1,2</sup>, Valentina Buneva<sup>1,2</sup>, Georgy Nevinsky<sup>1,2</sup>*

<sup>1</sup>*ICBFM SB RAS; <sup>2</sup>NSU, Novosibirsk, RUSSIA*

**Possibilities of enhancing the neuroprotective effect of autophagy activation in the brain by stimulation of an mTOR – independent pathway of its regulation in a transgenic mouse model of Parkinson's disease**

*Anna Akopyan, Aleksandr Pupyshev, Maria Tikhonova*

*Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

**Comparative analysis of the types of processing of visual information from the point of view of cognitive science**

*Alexandr Kashtnov, Mihail Pazhetnov, Elena Kashtanova*

*Novosibirsk State Technical University, Novosibirsk, Russia*

**Delta – and gamma – activity of resting state EEG as one of markers of risk of depressive disorders in migrants of subpolar and polar regions of Siberia**

*Natalya Milakhina<sup>1</sup>, Sergey Tamozhnikov<sup>2</sup>, Ekaterina Proshina<sup>2</sup>, Alexandra Karpova<sup>3</sup>, Alexander Savostyanov<sup>1</sup>, Elena Afanaseva<sup>3</sup>*

<sup>1</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>2</sup>*Institute of Physiology and Basic Medicine, Novosibirsk, Russia*

<sup>3</sup>*North – Eastern Federal University, Yakutsk, Russia*

**Reconstruction of Dementia Gene Network Using Online Bioinformatics Tools**

*Oleg Fateev<sup>1</sup>, Sergey Kovalev<sup>2,4</sup>, Yuriy Orlov<sup>3,4</sup>*

<sup>1</sup>*Institute of Pharmacy I.M. Sechenov First Moscow State Medical University, Moscow, Russia*

<sup>2</sup>*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*

<sup>3</sup>*Institute of Digital Medicine I.M. Sechenov First Moscow State Medical University, Moscow, Russia*

<sup>4</sup>*Novosibirsk State University, Novosibirsk, Russia*

**Interplay between 5 – HT and BDNF system in recombinant mouse strain upon chronic fluoxetine administration**

*Aleksandr Rodnyy, Elena Kondaurova, Yegor Antonov, Tatiana Ilchibaeva, Anton Tsybko, Vladimir Naumenko*

*Institute of Cytology and Genetics, SB RAS, Novosibirsk, Russia*