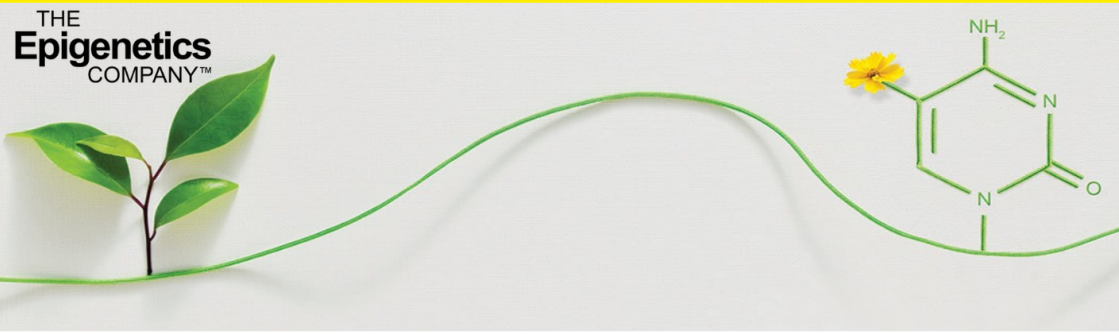
FEBS 2024
Advanced Course

Final programme

5th Danube Conference on Epigenetics

Research Centre for Natural Sciences, Budapest, Hungary

28-31 October 2024



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ACKNOWLEDGEMENTS

We would like to sincerely thank and express our appreciation to the following organisations for their support and assistance.

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Final programme

FEBS Advanced Lecture Course:
5th Danube Conference on Epigenetics
28-31 October 2024, Budapest, Hungary

Dear Colleagues and Friends,

We are delighted to welcome you to the **FEBS Advanced Lecture Course: 5th Danube Conference on Epigenetics** to be held between 28-31 October 2024, in the beautiful city of Budapest, Hungary. Building on the success of previous Danube Epigenetic meetings, we are thrilled to welcome you once again in person to Budapest.

The conference is meticulously organized by a distinguished team, including **Tamás Arányi** (RCNS / Semmelweis University, Budapest, Hungary), **Petra Hajkova** (MRC London Institute of Medical Sciences, London, UK), **Celia Martinez-Jimenez** (Helmholtz Pioneer Campus, Munich, Germany), **J. Andrew Pospisilik** (Van Andel Research Institute, MI, USA), **Lóránt Székvölgyi** (University of Debrecen, Debrecen, Hungary), and **László Tora** (IGBMC, Strasbourg, France).

Our primary goal is to foster collaboration among scientists in the fields of epigenetics, promoting intensive interdisciplinary interactions through this medium-sized meeting. The conference programme is strategically designed to offer numerous networking opportunities. We will have 150 participants including 24 outstanding invited lecturers.

We sincerely hope that you will find your participation attracting and useful. Wish you a pleasant event!

Tamás Arányi, MD, PhD

On behalf of the Organising Committee of 5th Danube Conference on Epigenetics

Scientific programme

Monday, 28 October, Day 1

10:00 – 18:30 *Registration*

Session 1: Data Visualization Workshop

chairs: Tamás Arányi and Lóránt Székvölgyi

10:15 – 11:00 **Cornelius Schneider**
Publishing Policies and Initiatives at EMBO Press

11:00 – 12:00 **Katarína Furmanová**
Pitfalls and Best Practices of Data Visualization

12:00 – 12:30 *Break*

12:30 – 14:00 **Jan Byška**
Data Visualization in Action: From Theory to Application

14:00 – 14:30 **Cecília Maria Arraiano**
FEBS Journals not only Publish Important Science but they also Promote Society: Examples from our Work in RNA, RNases and Control of Gene Expression

14:30 – 15:15 *Break*

s15:15 – 15:30 **Conference Opening**
chair: **Lóránt Székvölgyi**

Tamás Arányi
Opening Remarks

Session 2: Epigenetics and Development

chair: Ferenc Müller

15:30 – 16:00 **László Tora**
RNA Polymerase II Transcription is Supported by Partially Assembled TFIID Complexes in Embryonic Stem Cells

16:00 – 16:15 **Changqing Li**
The Structure of the Human TIP60/EP400 Super-complex: Merging Chromatin Remodeling and Histone Acetyltransferase Activities

16:15 – 16:30 **Inma Gonzalez**
MYC and MAX Drive the Reactivation of the Genome after Mitosis

16:30 – 17:00 *Coffee break*

Session 2: Epigenetics and Development
chair: Ferenc Müller

17:00 – 17:30 **Tineke Lenstra**
A Single-molecule Understanding of Transcriptional Bursting
EMBO Young INVESTIGATOR Lecture

17:30 – 17:45 **Acadia Kocher**
Probing the Chromatin and Location Dependencies of Transcription Factors in High Throughput

17:45 – 18:15 **Rob Klose**
Understanding How CpG Islands Regulate Gene Expression

18:15 – 18:30 *Technical break*

Session 3: Keynote Lecture
chair: Petra Hajkova

18:30 – 19:30 **Kenneth Zaret**
The Battle between Gene Activators and Silencers
IUBMB Keynote Lecturer

19:30 – 19:40 Gathering for the programme: in the lobby of the venue

20:00 – 22:00 ***Welcome reception***
ÖbölHáz Event Hall, 1117 Budapest, Kopaszi gát 1.
Reception is included in the registration fee

Tuesday, 29 October, Day 2

08:30 – 09:30	**Breakfast with the invited speakers** (maximum 3 students / speaker) **Registration for the programme is needed in advance**
09:00 – 10:00	*Registration*

Session 4: Epigenetics and Disease

chair: Tamás Arányi

10:00 – 10:30	**Michiel Vermeulen** Deciphering Lineage Specification During Early Embryogenesis using Multi-layered Proteomics
10:30 – 10:45	**Lehua Chen** TET1, Folate Resistance and Genetic Risk Factors in Neural Tube Defects
10:45 – 11:00	**Vivien Horváth** Local Heterochromatin Mitigates the Impact of a Transposable Element Insertion Causing a Neurodegenerative Disorder
11:00 – 11:30	*Coffee break*
11:30 – 11:45	**Mutay Aslan** All about FEBS
11:45 – 12:00	**Ferenc Müller**: Transcription Initiation Site Selection Defines Posttranscriptional mRNA Fate
12:00 – 12:15	**Ning Qing Liu** SMARCB1 Loss Activates Patient-specific Distal Oncogenic Enhancers in Malignant Rhabdoid Tumors
12:15 – 12:30	**Wayo Matsushima** Co-opted Retroviral Domain Defines Lineage-specific Enhancer Blocking Zinc-finger Proteins
12:30 – 13:00	**Kristian Helin** Epigenetics, and its Role in Transcriptional Regulation, Cell Fate and Cancer
13:00 – 14:30	*Lunch break*

14:30 – 16:00 **Poster session I.**

15:30 – 16:00 *Coffee break*

Session 5: Epigenetics and Differentiation

chair: László Tora

16:00 – 17:00 **Maria-Elena Torres-Padilla**
Epigenetic Mechanisms in Early Mammalian
EMBO Keynote Lecturer

17:00 – 17:15 **David Tremethick**
The Crucial Role of H2A.Z in Cell Fate Decisions: New Insights into the Regulation of Promoter Chromatin Architecture

17:15 – 17:30 **Sunil Kumar Pradhan**
Correlation of Epigenetic Dynamics with the Genome Replication Program in mESCs

17:30 – 17:45 *Apple break*

17:45 – 18:15 **Claire Rougeulle**
To Inactivate or Not: An eXISTential Question

18:15 – 18:30 **Vaidotas Stankevicius**
Chemical Epigenetic Approach for Specific Tracking of DNA Methylation in Mammalian Cells

18:30 – 18:45 **Gergely Nagy**
Lineage-determining Transcription Factor-driven Promoters Regulate Cell Type-specific Macrophage Gene Expression

18:45 – 19:15 **Petra Hajkova**
Stability, Turnover, and Erasure of Epigenetic Information

20:00 – *Beer session / Meet the lecturers*
Cactus Juice Gastro Pub, 1061 Budapest, Jókai tér 5.
Please note that the "Beer session" programme is not included in the registration fee, your consumption should be covered individually.

Wednesday, 30 October, Day 3

08:30 – 09:30 **Breakfast with the invited speakers**
(maximum 3 students / speaker)
Registration for the programme is needed in advance

Session 6: Epigenetics and Aging
chair: Andrew J. Pospisilik

10:00 – 10:30 **Celia Martinez-Jimenez**
Aging and Polyploidy: Dissecting a Novel Hepatocyte Cell State

10:30 – 10:45 **Yitzhak Reizel**
Hepatocyte Transplantation Remodels their Epigenetic Landscape and Reverses their Epigenetic Age

10:45 – 11:00 **Georgieva Milena**
Chromatin and Linker Histones: Their Role in Maintaining Genome Stability and Aging

11:00 – 11:30 *Coffee break*

11:30 – 12:00 **Andreas Beyer**
A Systems Perspective on Transcriptional Stress during Aging

12:00 – 12:15 **Margherita Mori**
Deciphering DNA Methylation Maintenance Mechanisms in Response to DNA Damage

12:15 – 12:30 **Epigenetic prize**

Ádám Sturm
N6-Methyladenine Dynamics in Nuclear and Mitochondrial Genomes: A Dual Pathway to Longevity

12:30 – 13:00 **Peter Tessarz**
Metabolism and Epigenetics in Aging

13:00 – 14:00 *Lunch break*

14:00 – 16:00 **Poster session II.**

15:30 – 16:00 *Coffee break*

Session 7: Keynote Lecture
chair: Celia Martinez-Jimenez

16:00 – 17:00 **Oded Rechavi**
Epigenetic Memories

Session 8: Epigenetics and Model Organisms
chair: Celia Martinez-Jimenez

17:00 – 17:30 **Eric Miska**
Non-coding RNAs in Early Development

17:30 – 17:45 **Ana Janeva**
Taking the HAT off: Perturbing Histone Acetylation for a Cell Fate Makeover

17:45 – 18:00 *Apple break*

18:00 – 18:30 **Ian Henderson**
Genetic and Epigenetic Regulation of the Arabidopsis Centromeres

18:30 – 18:45 **Richard Bartfai**
The Protein Landscape of the Chromatin States in P. falciparum Parasites

18:45 – 19:15 **Julie Ahringer**
Uncovering How the Genome Directs Development

20:30 - 22:30 **Gala dinner with river cruise**
Gala dinner is included in the registration fee

Thursday, 31 October, Day 4

08:30 – 09:30 **Breakfast with the invited speakers**
(maximum 3 students / speaker)
Registration for the programme is needed in advance

Session 9: Epigenetics and Inheritance
chair: Petra Hajkova

10:00 – 10: 30 **Nicola Iovino**
Epigenetics of Early Development
EMBO Young INVESTIGATOR Lecture

10:30 – 10:45 **Katalin Fejes Toth**
Beyond the Histone Code: an Unexpected Function of SetDB1 in Heterochromatin Spreading and Maintenance

10:45 – 11:00 **Archana Tomar**
Whispers of Diet: The Hidden Legacy of Sperm-borne Mitochondrial RNAs

11:00 – 11:30 *Coffee break*

11:30 – 12:00 **Susan Mango**
Dynamic Chromatin During Pluripotency and Gastrulation

12:00 – 12:15 **Juliane Glaser**
Enhancer Adoption by an LTR Retrotransposon Generates Viral-like Particles Causing a Developmental Limb Phenotype

12:15 – 12:30 **Christian Belton**
The Molecular and Phenotypic Features of Non-canonical Imprinting

12:30 – 13:00 **Andrew J. Pospisilik**
Probability, ERVs, and Phenotypic Variation

13:00 – 13:10 **Prizes and concluding remarks**

13:10 – 14:00 *Farewell snacks*

List of posters

P-01 **Agnieszka Belter**
The Epigenetic Effect of Cannabis on Glioblastoma Cell Lines

P-02 **Neethu Mohan**
Unraveling the Pharmacoepigenetic Effects of Antiepileptic Drugs

P-03 **Adina Aukenova**
Dimerization Controls Ikaros Transcription Factor Function

P-04 **Jure Krasic**
Cell-Free DNA Analysis of Blood and Ejaculate in Nonseminomatous Testicular Germ Cell Tumor Patients

P-05 **Péter Nánási**
Molecular Associations of Histone Variant H2A.Z Involving its C-terminal Tail

P-06 **Kristóf Gui**
The Analysis of the Classical and Extended Synergy Transcription Programs in Macrophages and in Epithelial-like Tumor Cells in the Presence and Absence of the Transcription Factor BACH1

P-07 **Teresa Rubio-Tomás**
Modulation of Life Span and Health Span upon Oxidative Stress by HDACs

P-08 **Nineļa Vainšeļbauma**
Polyploidy, Circadian clock Deregulation and Epigenetic Reprogramming to Reproductive Attractor States Promote Treatment Resistance in Malignant Tumors

P-09 **Logan Williams**
Cell-free DNA Methylation Patterns in Maternal Blood Plasma Show Overlapping Signatures Between Maternal Obesity and Child Autism Diagnosis, with a Substantial Enrichment for Genes Previously Implicated in Autism Susceptibility

P-10 **Julija Šmon**
Depression, Suicide and Proteinopathy: Elucidating the Relationships between Aggregation and Pathological Development (the DeSPERADo project)

P-11 **Chayeen Brotzki da Costa**
Investigating DNA Methylation Patterns During Annelid Regeneration

P-12 **Ibtissem Benhamza**
Epigenetic Modulation via the C-Terminal Tail OF H2A.Z

P-13 **Georgios Nikolopoulos**
Cockayne Syndrome B Protein is Implicated in Transcription and Associated Chromatin Dynamics in Homeostatic and Genotoxic Conditions

P-14 **Ksenia Kolobynina**
Novel Regulators of the Chromatin Response to DNA Damage

P-15 **Venkata Chelikani**
Regulation of Lactobacillus Acidophilus Gene Expression by Dietary Compounds via N4-Methylcytosine Modifications: A Novel Approach to Probiotic Engineering

P-16 **Esteban Retamales**
Intergenerational Adaptation of C. elegans to High Temperature Induced Diapause

P-17 **Priyanka Mehra**
MicroRNA Sequencing Reveals New Targets to Understand the Pathology of Fat Accumulation in Duchenne Muscular Dystrophy

P-18 **Leonard Steg**
Exploring Mouse Strain Specific Effects of Low Protein Diet on Liver Epigenome and Transcriptome and on Serum Lipidome

P-19 **Teodora Lukic**
Unraveling ΔNp63-Driven Transcriptional Reprogramming in Squamous Carcinogenesis: Insights from Comparative Omics Analysis

P-20 **Maria Teresa Alejo Vinogradova**
Investigating PRC1 Variants Dynamics during Mouse Embryo Development

P-21 **Kajsa Karlsson**
Small RNAs in Sperm – Paternal Contribution to IVF Outcomes

P-22 **Gábor Pápai**
Binding to Nucleosome Poises Human SIRT6 for Histone H3 Deacetylation

P-24 **Constance Merdrignac**
Autism-related gene Intergenerationally Regulates Neurodevelopment and Behavior in Fish through Non-genetic Mechanisms

P-25 **Emma Torun**
Investigating the Role of the Deubiquitinase BAP1 at Enhancer Regions

P-26 **Alice Jouneau**
Changes in H3K4me3 and H3K27me3 Landscapes During in vitro Derivation of Embryonic and Trophoblast Stem Cells – a Bovine Study

P-27 **Zsófia B. Nagy**
Assessment of the Environmental Factors Affecting DNA Stability on Leukocytes

P-28 **Kristine Salmina**
Mitotic Slippage, Soma-germ and Ameboid Transitions Cooperatively Support Tumour Cell Resistance after Genotoxic Treatment

P-29 **Barbara Bartak**
Examination of Epigenetic Changes in Patients with Hyperhomocysteinemia after Folic Acid Supplementation

P-30 **Ferenc Jankovics**
Identification of Structural Elements of the Sov Protein Involved In Heterochromatin-Mediated Transcriptional Transposon Silencing in Drosophila

P-31 **Inge de Krijger**
Uncovering Enhancer Dynamics in B-cell to Macrophage Transdifferentiation using Proximity Proteomics

P-32 **Hiba Souaifan**
Role of HP1 Proteins in the Maturation and Degradation of Non-coding RNAs in Connection with the Nuclear RNA Exosome Complex

P-33 **Ehsan Pourkarimi**
High Tryptophan Concentration Leads to Genomic Instability, Changes in Chromatin Structure, and Gene Expression in *Caenorhabditis elegans*

P-34 **Amélie Taschereau**
Placental microRNA are Associated with Insulin Secretion During Pregnancy

P-35 **Manu Araolaza**
Chronic Morphine Exposure Enhances Pluripotency in mESCs Through DNA Methylation

P-36 **Bela Vasileva**
Could Phytometabolites be the Next Breakthrough in Anti-ageing Therapy?

P-37 **Dávid Kis**
Ultrasensitive Early Detection of Colorectal Cancer by Cell-Free DNA Methylation-based Liquid Biopsy Assay

P-38 **Tamás Linkner**
Characterization of Embryonic and Mesenchymal Properties of Tumor Cells with Flow Cytometry

P-39 **Péter Hunyadi**
Inferring DNA Methylation from Low-pass WGS of Cell-free DNA using Fragmentation Patterns in Ovarian Cancer Samples

P-40 **Mirabela Romanescu**
Characterisation of the hsa-miR-130a and hsa-miR-365a -3p/-5p Ratios in Prostate Cancer Biospecimens

P-41 **Aimée Rodica Chis**
MIR-19 as Levodopa Therapy Responder in Parkinson's Disease

P-42 **Paula Diana Ciordas**
Characterization of Plasma hsa-miR-19b-3p as Prognostic Biomarker in Non Small Cell Lung Cancer

P-43 **Douwe ten Bulte**
Understanding the Role of DNA Methylation at H3K9me3-decorated Heterochromatin

P-44 **Ovidiu Sirbu**
Consistent Discordances in Quantification of 3p and 5p microRNA Species are Independent of their GC Content and Secondary Structure Stability

P-45 **Laura Quispe**
Experimental Design to Study Sperm DNA Methylation in Response to Elevated Temperature in Medaka (Oryzias latipes) and its Transgenerational Transmission

P-46 **Amelie Bonnet-Garnier**
Abnormal Organization of Major Satellite Sequences in Zygote of Mice Knock-out for Suv39h1-Repressive Antisense lncRNA

P-47 **Teruhito Ishihara**
Nanosplit - Allele-specific Read Sorting Programme for Oxford Nanopore long Read Sequencing Platform

P-48 **Alexandra Mocanu Dobranici**
Highlighting the role of Linker of Cytoskeleton and Nucleoskeleton (LINC) Complex in Magnetically-driven Epiegentic Changes

P-49 **Máté Varga**
Zebrafish Models of Defective Pseudouridylation Provide Insights into the Ethiology of Ribosomopathies

P-50 **Damir Baranasic**
Functional Impacts of Promoter Evolution in Cyprinus Carpio Carpio Revealed by High-resolution CAGE Analysis

P-51 **Sára Zsigrai**
Folic Acid and S-Adenosylmethionine are Effective Methylome Modulators in Colorectal Cancer Cell Lines

P-52 **Kristóf Rada**
Long-read Nanopore Sequencing in Pursuit of Complete Methylation Profiling of Hungarian CRC Patients

P-53 **Krisztina Szigeti**
Global DNA Hypomethylation Linked to Decreased Methyl-donor Level in Colorectal Cancer Progression

P-54 **Eszter Farkas**
Effects of Folic Acid Levels on DNA Fragmentation and Epigenetic Changes

P-55 **Fruzsina Bányai**
Monitoring Epigenetic Changes after Folic Acid Supplementation

P-56 **Eleftheria Chatzantonaki**
The Functional Role of Polycomb-mediated Chromatin Architecture During Neuronal Development.

P-57 **Firuze Unlu Bektas**
Investigating MBD5 and its Contribution to PR-DUB Complex Function in the Developing Brain

P-58 **Kian Peng Koh**
The Epigenetic Functions of Vitamin C, TET1 and Genetic Susceptibility for Neural Tube Defects.

P-59 **Liepa Gasiule**
Metabolic Labeling of Individual DNA Methylomes in Live Cells

P-60 **Antonio Adamo**
YMETHX: Deciphering the Interplay of DNA Methylation and Gene Expression in Male Sex Chromosome Aneuploidies

P-61 **Ankan Ankan Roy**
Hub Genes in Driving Colon Cancer Progression: Plasma Membrane Lipid Raft Signaling Regulates The Hub Genes by Epigenetic Modulation

P-62 **Tamás Arányi**
Improve-RRBS: a Novel Tool to Correct the 3′ Trimming of Reduced Representation Sequencing Reads

P-63 **Zsuzsanna Gaál**
Targeted Epigenetic Interventions in Cancer with an Emphasis on Pediatric Malignancies

P-64 **Tamás Arányi**
Hepatocyte-Specific DNMT3A And DNMT3B Knockout Mice Develop Pre-Cancerous Phenotype

P-65 **Anna Poscher**
Synergistic Interaction Between IL-4 and Hypoxia: An Epigenetic Perspective

P-66 **Dániel Tóth**
Involvment of de novo DNA Methyltransferases in Stress-Induced Responses of the Liver

P-67 **Ningqing Liu**
Targeting Transcription Factor Addiction to Treat Acute Myeloid Leukemia

P-68 **Zsófia Varga**
Identification of Synergistic Interactions Between the Opposing Polarization Signals in Macrophages

P-69 **Tomasz Wojdacz**
Blood cells of BRCA1 Mutation and Epimutation Carriers Appear to Acquire Specific Epigenetic Signatures

P-70 **Muhyiddeen Muazu**
Postnatal Hyperthyroidism Causes DNA Hypermethylation and Marked Decrease of Dio1 Gene Expression in the Liver of Adult Mice

P-71 **Jan Bińkowski**
Blood and Urinary Arsenic Levels are Associated with Methylation of Promoters of Genes Involved in Molecular Processes Related to Arsenic Toxicity

P-72 **Sarah Offley**
TRanscriptional Regulation by Non-canonical PP2A Complexes

P-73 **Katarzyna Sokołowska**
Methylation Levels at Non promoter CpG sites are Correlated with Expression of Distant Genes

P-74 **Krystyna Giemza**
Precision Chromosome Engineering to Study Nucleolar Formation in Human Cells

P-75 **Ádám Sturm**
N6-Methyladenine Dynamics in Nuclear and Mitochondrial Genomes: A Dual Pathway to Longevity

P-76 **Lenka Steiner Mrázová**
Classification of Variants of Uncertain Significance in Neurological Diseases: Three Different Approaches

P-77 **Agnieszka Zelisko-Schmidt**
Efficient, Low-input Chromatin Profiling using iDeal CUT&Tag and ATAC-seq Kits

P-78 **Tibor Csorba**
Transcriptional Control of Heat Stress Response in Plants

General information

Venue

Research Centre for Natural Sciences (RCNS)
Budapest
Magyar Tudósok körútja 2.
H-1117

Opening hour of the registration desk

Monday, 28 October	10:00-19:00
Tuesday, 29 October	09:00-19:00
Wednesday, 30 October	09:30-19:00
Thursday, 31 October	09:30-13:00

Important phone numbers

English is usually spoken at the emergency numbers listed below.
Central help number: 112

Ambulance: 104
Fire brigade: 105
Police: 107
Central help number: 112
General enquiries: 197
Inland enquiries: 198
International enquiries: 199
Hungarian Automobile Club help number: 188

Social programmes

Welcome reception

Monday, 28 October 20:00-22:00
ÖbölHáz Event Hall (1117 Budapest, Kopaszi gát 1.)

Gathering for the programme: in the lobby of the venue at 19:40.
The organisers invite you for a dinner on the evening of your arrival. We wish to serve you some refreshment and a buffet dinner after your travel. Our other aim is to create a familiar atmosphere where you can meet old friends, and make new relationships, too.
Included in the registration fees. Entry by ticket only.

Beer session

Tuesday, 29 October, from 20:00
Cactus Juice; 1061 Budapest, Jókai tér 5.

On the second day of the conference we are planning to have a "Beer session" in a local pub during the evening. We would like to invite you to join us that night, to have the opportunity to get to know each other better.
Please note that the "Beer session" programme is not included in the registration fee, your consumption should be covered individually.
How to get there: at Petőfi híd (Buda side) tram stop, take tram line number 4 or 6 to the direction of Széll Kálmán tér and get off at Oktogon stop (this is the ninth stop). The pub is a two-minute walk from Oktogon.
Optional programme.

Gala dinner with river cruise

Wednesday, 30 October 20:30-22:30

Come and enjoy a friendly and memorable gala dinner with the conference participants on an amazing river cruise. A delicious dinner with wines will be served during the night. When in Budapest one cannot miss out on sailing past the architectural wonders of the Hungarian Capital. Our gala dinner with river cruise gives you the opportunity to admire in the breath-taking panorama of the capital. **Please note: there are no stops along the cruise route.**

The boat can be boarded on the other side of the Danube (Pest side), at the Boráros tér pier. Please make your way to the pier independently by 20:15 at the latest.
The boat will depart at 20:30 and will return to the same place at 22:30.
Individual boarding is not allowed!
Included in the registration fees. Entry by ticket only.

Presenters' guidelines

Invited presentations
Each presentation takes 20 minutes +10 minutes discussion.

Oral presentations
Each presentation takes 10 minutes +5 minutes discussion.

Technical instructions:
Please prepare your presentation in .ppt, .pptx or .pdf file. If you wish to have a video, please contact the technician in the lecture hall during the break before your presentation (or preferably earlier) to check it in advance. Please note that using your own notebook is not recommended. **Please upload your presentation at latest during the break before your session.**

Posters

Posters will be mounted during the whole Conference. Please make sure to remove your poster till the given deadline.

Mounting: 28 October, from 12:00
Removal: 31 October, before 12:00

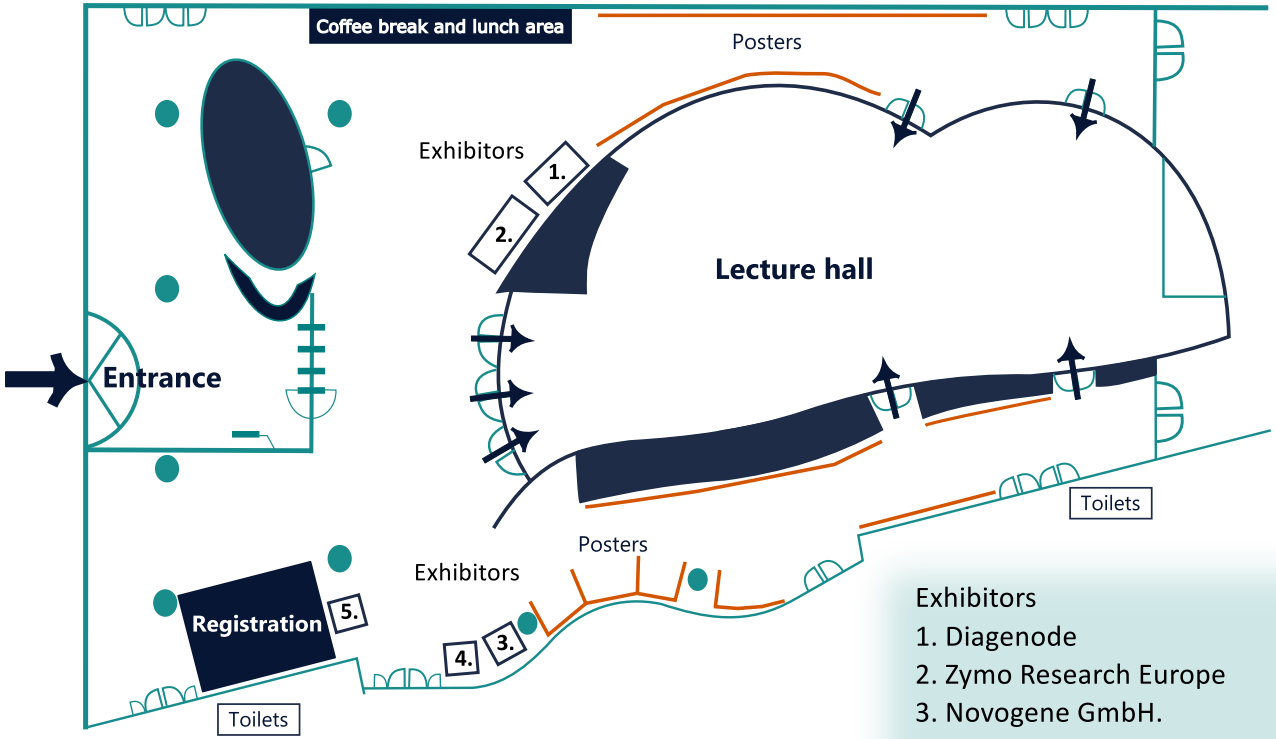
Conference mobile application

5th Danube Conference on Epigenetics is supported by the SmartEvents mobile application. Using the application is free for all registered delegates and the content used therein is accessible only for our participants. The application helps you to find all information regarding the event, manage your daily agenda, download materials, communicate with all the other delegates on the event and more. All changes of the programme (if any) will be tracked in the device in real time, and the whole conference will be IN YOUR HAND all the time. For further help, please contact our colleagues at the registration desk.

FEBS Advanced Lecture Course:
5th Danube Conference on Epigenetics

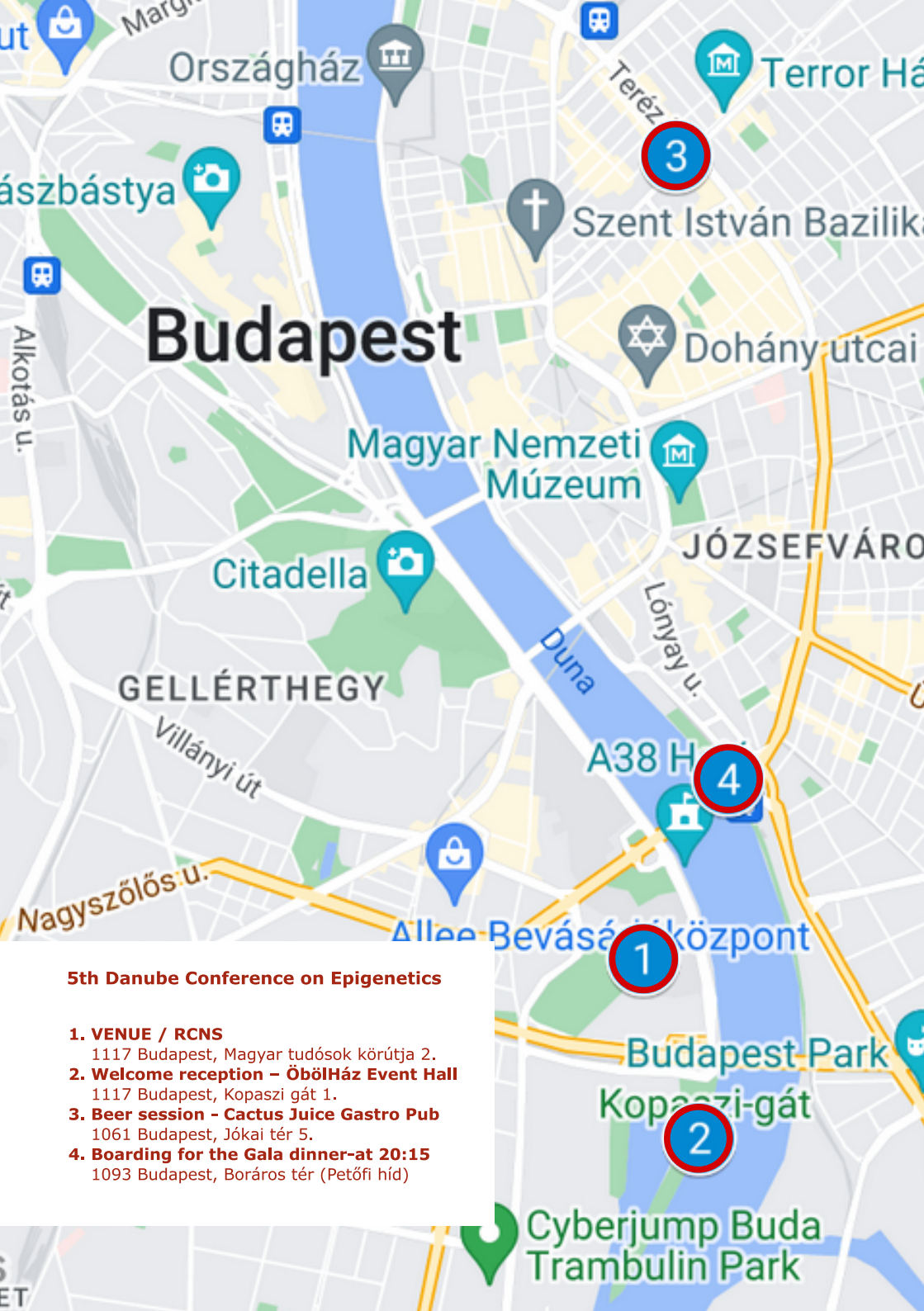
28-31 October 2024

Research Centre for Natural Sciences, Budapest, Hungary



Exhibitors

1. Diagenode
2. Zymo Research Europe
3. Novogene GmbH.
4. HaploX GeneTech (Hong Kong)
5. Merck Life Science Ltd.



5th Danube Conference on Epigenetics

1. **VENUE / RCNS**
   1117 Budapest, Magyar tudósok körútja 2.
2. **Welcome reception – ÖbölHáz Event Hall**
   1117 Budapest, Kopaszi gát 1.
3. **Beer session - Cactus Juice Gastro Pub**
   1061 Budapest, Jókai tér 5.
4. **Boarding for the Gala dinner-at 20:15**
   1093 Budapest, Boráros tér (Petőfi híd)

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