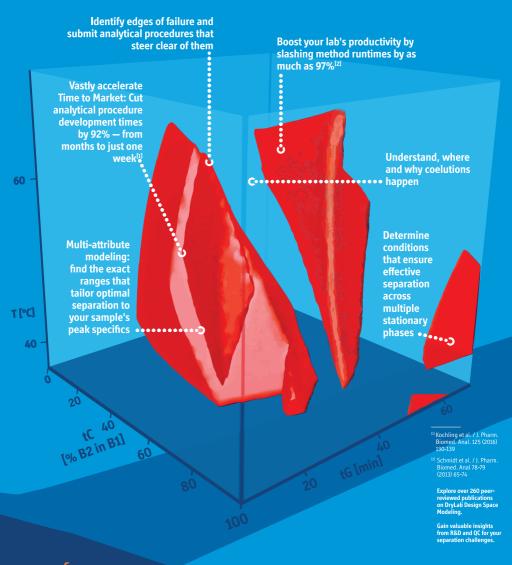


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GENERAL INFORMATION

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Magyar Biokémiai Egyesület (Hungarian Biochemical Society)

ORGANISING COMMITTEE

László Buday¹, Mihály Kovács², Beáta Lontay³, Zoltán Lipinszki⁴, András Micsonai², Virág Vas²

- HUN-REN Research Centre for Natural Sciences Magyar tudósok körútja 2., Budapest, H-1117 https://www.ttk.hun-ren.hu/ei/en/
- 2 Eötvös Loránd University Pázmány Péter sétány 1/c., Budapest, H-1117 https://www.elte.hu/
- Department of Medical Chemistry, University of Debrecen Egyetem tér 1., Debrecen, H-4032 https://chemistry.med.unideb.hu/en
- 4 FEBS Member Society Representative University of Szeged Dugonics tér 13., Szeged, H-6720 https://u-szeged.hu/english

29-31 August, 2024, Budapest, Hungary



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https://hbs-conference.hu

WEBSITE OF THE HUNGARIAN BIOCHEMICAL SOCIETY

http://www.mbkegy.hu

VENUE

Campus of Eötvös Loránd University

1117 Budapest, Pázmány Péter sétány 1/A

OPENING HOURS OF THE REGISTRATION

Thursday, 29 August, 2024	-	12.00 – 18.00
Friday, 30 August, 2024		08.30 - 18.00
Saturday, 31 August, 2024		08.30 - 13.30

ONSITE CONTACT NUMBERS

Ildikó Gémes-Kruppa / Attila Varga

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OFFICIAL LANGUAGE

Official language of the Conference is English (no translation is available).

REGISTRATION FEES

Registration fees include admission to the scientific sessions and to the exhibition, Welcome reception, Coffee breaks, Lunches, VAT.

Tickets for the gala dinner can be requested through the registration database. Participation in the dinner is possible upon payment of 15.000 HUF.



FXHIBITON

In accordance with the conventions of the conference, parallel to the scientific sessions a professional exhibition is to be organised. Please have a look at the exhibition floor plan of the booklet.

SOCIAL PROGRAMMES (incl. in the registration fees)

Thursday, 29 August, 2024 Welcome reception (Venue)

Friday, **30 August**, **2024** Gala dinner (Öböl Ház) – participation only with ticket

Friday - Saturday, **30-31 August 2024** Lunches (Venue)
Thursday - Saturday, **29-31 August 2024** Coffee breaks (Venue)

(indicated in the programme)

All participants and accompanying persons will receive a personal badge upon registration. You are kindly requested to wear your name badge when attending the meetings or social events.

CANCELLATION POLICY

Cancellations on registration can be made only in writing. The refund for cancellations made prior to 8 July, is 100%. After this there is no way to refund in case of latter cancellation.

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The price of the ordered services will be indicated on the final invoice according to the Hungarian official financial rules. Official final invoices and receipts for fees paid by the participants are sent by email right after the payment arrives to our bank account. Please forward them to the financial department of your institution.

LIABILITY AND INSURANCE

The organisers cannot accept liability for any personal accidents, loss of belongings or damage to private property of participants and accompanying persons that may occur during the conference.

POSTER PRESENTATION SCHEDULE

Posters should be affixed and removed by the presenters themselves in time. All posters will be on display throughout the whole conference from P-01 to P-68. Posters should be mounted from the time of registration, on 29 August and they have to be removed latest during the last coffee break till 11 AM on 31 August.

ODD numbered posters will be presented on Thursday (Poster discussion I.) between 17:00-18:00, EVEN numbered posters will be presented on Friday (Poster discussion II.) between 16:45-18:00.

29-31 August, 2024, Budapest, Hungary



PROGRAMME



Day 1 - Thursday, 29 August 2024

15:00-15:10 *Opening*

15:10-16:20 Plenary session

Congress Hall
Chair: László Buday

15:10-16:00 PL-1

György Keserű

Mapping protein-protein interaction surfaces by photoactivable molecular fragments

16:00-16:20

Bio Science award talk

Áron Szabó

LC3-associated phagocytosis promotes glial degradation of axon debris after injury in Drosophila models

16:20-16:35 Company lecture of Kromat Kft.

Éva Csősz

Complex analysis of obesity and type 2 diabetes moonlights Olink's proximity extension assay as a method of choice in multiomics setup

16:35-16:50 Company lecture of Molnar Institute

Imre Molnár

Speeding up research in Biochemistry by Modeling HPLC

16:50-17:15 *Coffee break*

17:00-18:15 Poster discussion I. – Odd numbers

18:15- Welcome reception – Gömb Aula of the Venue

29-31 August, 2024, Budapest, Hungary



Day 2 - Friday, 30 August 2024

09:00-09:30	Plenary session Congress Hall Chair: László Virág
09:00-09:30	PL-2 József Tőzsér Structure, function, inhibition and resistance development of sars-cov-2 main protease (MPRO). what is common with retroviral proteases?
09:30-10:50	Molecular signaling, cell-cell communication, cell death and differentiation Chair: Zoltán Lipinszki

O-01 – Zoltán Lipinszki

Exploring the Substrate Recognition Mechanism of Protein Phosphatase 4 Reveals Novel Functions of the Enzyme in DNA Damage Repair

O-02 - Krisztina Tar

Can Manipulating Mitochondrial Dynamics Enhance Neuronal Differentiation?

O-03 - Endre Kristóf

Upregulation of Inhibitor of DNA Binding 1 and 3 is important for efficient thermogeninc response in human adipocytes

O-04 - Virág Vas

A comprehensive study from EMT regulation to cancer biomarker discovery: "Tyrosine kinase substrate with four SH3 domains" in focus

10:50-11:10 *Coffee break*



11:10-12:30

Regulation of gene expression, regulatory RNA, epigenetics Chair: Mihály Kovács

O-05 – Mihály Kovács

Subcellular rearrangements of bacterial single-stranded DNA binding (SSB) protein condensates indicate roles in stress response and cell proliferation

O-06 - Ágnes Tantos

Competing endogenous RNA networks in different cancers

O-07 - Pantazi Vasiliki

K63-linked ubiquitylation of RNAPII in response to DNA Double-Strand Breaks drives repair regulation

O-08 - Jacksi Mevan

The role of EZH2 in TKS4-deficient colorectal carcinoma cells.

12:30-13:30 Lunch

13:30-14:50

Protein structure, function and modeling, proteomics Chair: Ferenc Erdődi

O-09 - Orsolya Tőke

Pressure denaturation of proteins

O-10 -Zoltán Kónya

Characterization and analysis of protein phosphatase-1 and -2A modulators identified by high throughput screening

O-11 – Boglárka Zámbó

Proteome-wide affinity survey of DNA-protein interactions: A case study on the erythroid-specific promoter of ATP2B4 gene and GATA1 mutations

O-12 - Gergő Gógl

The non catalytic epsilon DNA polymerase subunit is an NPF motif recognition protein

14:50-15:10 *Coffee break*

29-31 August, 2024, Budapest, Hungary



15:10-16:3	

Bioinformatics, synthetic biology, genome engineering, biotechnology Chair: Zoltán Ivics

O-13 - Zoltán Ivics

Advances in non-viral engineering of therapeutic cells with the Sleeping Beauty transposon system

O-14 - Orsolya Kapuy

Re-wiring the AMPK-ULK1 regulatory connection at the service of periodic autophagy induction

O-15 - Regina Martinek

Comprehensive investigation of the mutagenic potential of six pesticides classified by IARC as probably carcinogenic to humans

O-16 - István Hajdú

Beta secretase 1 Recruits Amyloid beta Precursor Protein to ROCK2 Kinase, Resulting in Erroneous Phosphorylation and Beta¬amyloid Plaque Formation

	Chair: Virag Vas
16:45-18:00	Poster discussion II. – Even numbers

18:30 Gala dinner – Öböl Ház, participation only with ticket



Day 3 - Saturday, 31 August 2024

09:00-10:40

Molecular basis of disease and therapy, stem cells, immunity and inflammation

Chair: Zsuzsanna Izsvák

O-17 – Zsuzsanna Izsvák

HERVHLin protects human genome from mutagenic retroelements by rewiring the conserved LIN28A/let7 pathway in the early embryo

O-18 - Flóra Szeri

Alkaline phosphatase activity correlates with circulatory mineralization factors upon aging in mice and human

O-19 - Rini Arianti

Solute carrier transporters-mediated amino acids uptake is essential for thermogenic activation of human primary adipocytes

O-20 - Andrea Kocsis

SARS-CoV-2 Nucleocapsid Protein Is Not Responsible for the Over-Activation of Complement Lectin Pathway

O-21 - Yousif Qais Muhsin Al-Khafaji

Short-term thiamine deficiency altered transcriptomic profile of human deep cervical-derived adipocytes

10:40-11:00 *Coffee break*

29-31 August, 2024, Budapest, Hungary



11:00-12:20

Translational research, medicinal biochemistry, drug development Chair: György Vereb

O-22 – György Vereb

Chimeric antigen receptors (CAR) in the treatment of solid tumors: lessons from HER2+ preclinical models

O-23 - József Prechl

A truly quantitative antibody serological assay utilizing double-titration on antigen microspots

O-24 - Mohammed Maki

Anaerobic glycolysis and MCT1inhibition via diclofenac lead to tumorigenesis regression, ROS accumulation through links intracellular acidification and OXPH in K562 erythroleukemia

O-25 - Tibor Pankotai

Proteogenomics reveals varying levels of intra-patient heterogeneity in melanoma metastases

12:20	Closing ceremony
12:40-	Lunch



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List of posters

P-01 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-02 Botond Engel

The mutagenic outcome of translesion replication at DNA adducts depends on polymerase choice

P-03 Valentina Madár

Exploring multicellular growth of yeast

P-04 Judit Tóth

dUTPase modulates mycobacterial homologous recombination through interaction with AdnAB

P-05 Ankita Murmu

Intercellular communication and CD8+ T-cell trajectory analysis in ovarian cancer tissues

P-06 Anna Zichy

Mapping and validation of structural variants in lung cancer for enhanced understanding of tumor development

P-07 Tamás Takács

Orthosteric inhibitors targeting SHP2: structural insights and challenges

P-08 Eszter Vámos

The effect of trimethylamine, a metabolite of gut microbiota on the metabolic reprogramming of M1 macrophages

P-09 Nikolett Nagy

Moonlighting role for dUTPase: establishing a viable mammalian cell line with high uracil DNA levels

P-10 Bence Farkas

The dominant form of the C1 inhibitor R444C variant causing type II hereditary angioedema is its covalent adduct with serum albumin explaining elevated levels

P-11 András Zeke

Analyzing redox features of proteins based on AlphaFold2 models

P-13 Péter Tóth

Cell sorting based method for identifying CRISPR knock-in clones

P-14 Kriszinta Rubina Vass

Recombinant production and characterization of human alpha-keto acid dehydrogenase complex subunits and their disease-relevant variants

P-15 Balázs Csoma

Prime editing: advancing the frontiers of genetic engineering

P-16 Ilka Keller

Smoothelin-like protein 1 promotes insulin sensitivity and modulates the contractile properties of endometrial epithelial cells with insulin resistance

P-17 Zsuzsanna Biczók

A study on the reduction of self-complementarity in the prime editing RNA

P-18 Károly Hegedűs

Estimating efficiency of immunosuppressive therapy in heart transplant patients

P-19 Bálint Csörgő

Repurposing CRISPR-Cas3 systems for probing phage-host interactions

P-20 Eszter Holub

Key factors and consequences of drug induced genomic cytosine deamination

29-31 August, 2024, Budapest, Hungary



P-21 Balázs Ligeti

ProkBERT: genomic language models for understanding the microbiome

P-22 Alexandra Sándor

Development of an induced pluripotent stem cell-based organoid model for investigating the molecular mechanisms of cerebellar ataxia

P-23 Csaba Kiss

Investigating the role of non-coding genetic variants in colorectal cancer through single-cell multiomics analysis

P-24 Edit Szabó

Genetic variants of the human thiamine transporter (SLC19A3, THTR2) - potential relevance in type 2 diabetes mellitus

P-25 **Ágnes Becsei**

Detecting uracil in DNA with nanopore sequencing

P-26 Katalin Vincze

Exploring the neurological impact of type 2 diabetes in induced pluripotent stem cell derived neuronal progenitors and neurons

P-27 **Beáta Vértessy**

Artificial Intelligence-based search for new enzymes to degrade polluting hydrocarbons

P-28 **Hédi Maczelka**

Developing a novel endothelial dysfunction model using pluripotent stem cell-derived endothelial cells for extracellular matrix recellularisation

P-29 Dániel Molnár

Antibacterial action of membrane-incising peptide lamellae

P-30 Zita Bognár

Effect of Desethylamiodarone on mitochondrial mechanisms in breast cancer cell lines

P-31 Dávid Kovács

Triglyceride acyl length drives lipolysis and adipocyte metabolism

P-32 **Ádám Ungvári**

Myosin phosphatase promotes insulin-senitizing mechanisms in C2C12 muscle cells

P-33 Mihálv lózsi

Functional characterization of complement Factor H CCP19-20 variants with a set of host-derived and microbial ligands

P-34 Viktória Jenei

Immunological characterization the different forms of necrotic cell death

P-35 János András Mótyán

Structure-based identification of interacting partners of the human retroviral-like aspartic protease 1 (ASPRV1)

P-36 Renáta Bíró

ARHGAP25 has an anti-inflammatory role in the imiquimod-induced psoriasis mouse model

P-37 Kiarie Irene Wanjiru

Inhibition profiling of integrase strand-transfer inhibitors and the capsid inhibitor against HIV-2, and exploration of factors associated with viral latency

P-38 Bálint Jezsó

Phase separation and cytoplasmic granulation by single-stranded DNA binding protein 1 (SSB1) delineate new mechanisms of cellular stress response



P-39 **Éva Gráczer**

Dissecting the cellular effects of beta-actin mutations causing non-muscle actinopathies

P-40 Zoltán Ódor

Utilizing NanoFlare gold nanoparticles for dynamic monitoring of gene expression and cellular differentiation in colorectal cancer cells

P-41 Mária Golda

Investigation of the substrate specificity of human retroviral-like aspartic protease 1 (ASPRV1)

P-42 Viktória Tisza

CHD1 deletion in prostate cancer and its impact on PARP inhibitor sensitivity

P-43 Kriszinta Rubina Vass

NMR structural analysis of the lipoyl domain of the hKGDHc-E2 protein

P-44 Petra Körmendi

Crosstalk of LKB1- and PKA-dependent signaling pathways in the regulation of hepatocyte polarity

P-45 Dávid Szakács

Interaction of complement C3b with factor H family proteins

P-46 Seo Mizuki

Transferrin receptor-mediated iron uptake is required for thermogenic activation in human cervical derived adipocytes

P-47 Dávid Szakács

Annexin A2 is a ligand for complement factor H, FHL-1 and FHR proteins

P-48 Mustafa Abdulkareem

Investigating the role of RNA binding in the cellular function of KMT2D

P-49 Mónika Ágnes Szütsné Tóth

Structure-function coordination in the SALS protein

P-50 Otília Tóth

Expression of dUTPase in mouse postnatal development

P-51 Péter Gaszler

Structural analysis of the EB1 protein

P-52 Tamás Orbán

Exosomal small RNA profiling in first-trimester maternal blood explores early molecular pathways of preterm preeclampsia

P-53 Mohamad Mahfood

Expression of myosin phosphatase in breast cancer

P-54 Viktória Perey-Simon

Role and effect of uracil metabolism on zebrafish embryonic development

P-55 Csaba Magyar

MD simulation analysis of heterodimeric mutual synergistic folding proteins

P-56 Laura Vizkeleti

Functional understanding Septin9 intronic DNA methylation and its impact on colorectal cancer development and diagnosis

P-57 Rita Hirmondó

Do Mycobacteria really have no actin? Characterization of a potential Mbl protein

P-58 Tamás Arányi

Hepatocyte-specific DNMT3A and DNMT3B knockout mice develop pre-cancerous phenotype

P-59 Zoé.S. Tóth

The full-length Stl repressor is required for efficient inhibition of mycobacterial dUTPase

29-31 August, 2024, Budapest, Hungary



P-60 **Zsuzsánna Réthi-Nagy**

Correlation between virulence and cytotoxicity of multidrug-resistant bacteria in human cell models

P-61 Katalin Kuffa

Side-specific cell surface biotinylation to analyze the apical-basolateral distribution and accessible regions of transmembrane proteins in polarized epithelial cells

P-62 Richárd Kinter

The effect of fluorescent photoactive Hsp90 inhibitors on the metastatic properties of melanoma cells

P-63 Anna Sánta

Analysis of disease-associated point mutations of the Shank1 PDZ domain

P-64 Doa'a Al-u'datt

Nanoparticles-derived royal jelly exerts antidiabetic properties

P-65 Eszter Nagy-Kanta

Structural investigation of the multivalent complex of the GKAP scaffold and LC8 hub protein

P-66 Nikoletta Pfeiffer

Effect of olaparib on temozolomide-induced cell death

P-67 Bálint Péterfia

Ensemble-based modeling using SAXS and NMR data of the Drebrin single alpha-helix

P-68 Ammar Darwish

Silver-lipid nanoparticles shaped for targeted tumor therapy

P-69 Mátyás Sámuel Szabó

Examining the molecular mechanisms of irradiation-induced myocardial damage

P-70 Barbara Erdélyi-Furka

The cytoprotective effect of preimplantation factor on irradiation-induced cardiac cell injury

P-71 **Dóra Nógrádi-Halmi**

Kynurenic acid protects cardiac cells against doxorubicin-induced cardiotoxicity

P-72 Renáta Gáspár

Mechanism of kynurenic acid-derived protection against simulated ischemia/reperfusion-induced cardiac cell damage: investigation on the potential modulation of autophagy

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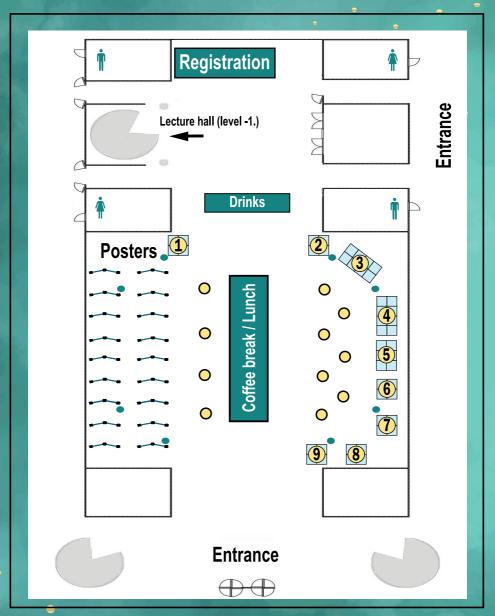


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