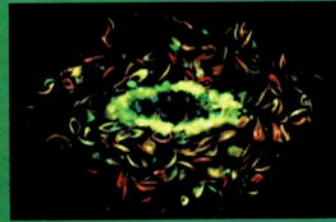
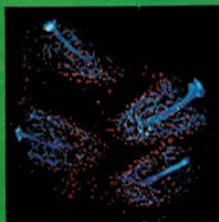
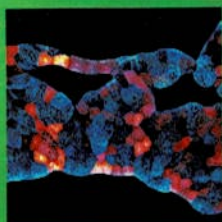
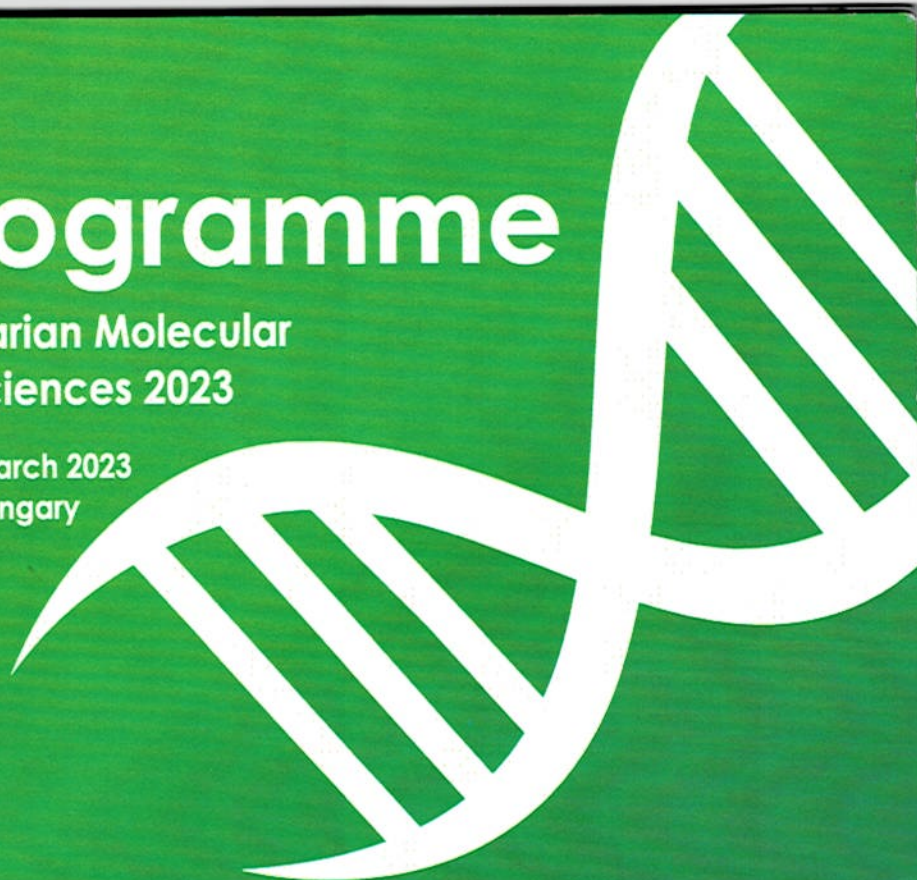


Programme

Hungarian Molecular
Life Sciences 2023

24-26 March 2023
Eger, Hungary

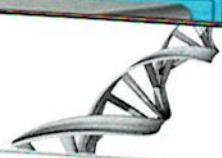


Hungarian
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Hungarian Genetics Society



List of posters - 24-25 March 2023

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| P-001 | Dóra Halmi
Mechanism of cytoprotection by kynurenic acid against simulated ischemia/reoxygenation-induced cardiac cell damage: a focused investigation on autophagy |
| P-002 | Marianna Holczer
The study of autophagy-dependent survival in endoplasmic reticulum stress |
| P-003 | Fanni Keresztes
Autophagy-related small GTPase proteins' effect on <i>Drosophila</i> lifespan and Parkinson's disease model |
| P-004 | Zoltán Köhler
Effect of tilorone on glucose homeostasis in a high-fat diet mice model |
| P-005 | Dorina Lenzinger
MVB-like extracellular vesicles |
| P-006 | Győző Szenci
The role of Ankyrin-repeat palmitoyl-transferases in lysosomal degradation |
| P-007 | Szabolcs Takáts
Uncovering degradation independent functions of autophagy in cancer |
| P-008 | Maher Alnajjar
A large family-based approach for determining the parent's haplotype, recombination events, de novo mutations, and sequencing errors detection |
| P-009 | Ákos Dörmő
Comparison of technical approaches applied in gut microbiome analyses reveals significant inconsistencies |
| P-010 | Tibor Hajdú
Proteomic and network biology analysis of the plasma membrane proteome in human cutaneous melanocytes and melanoma cells |
| P-011 | Márk Kovács-Valasek
Identification of novel cryptic primycin biosynthetic gene clusters within the pseudonocardiaaceae family |
| P-012 | Balázs Liget, Zsófia Molnár
Transfer learning for microbiome based predictive disease classification |
| P-013 | Kolos Nemes
Predicting drug response in small cell lung cancer cell lines using gene expression signatures |
| P-014 | Erda Qorri
Decoding canine senescence: A single-cell RNA-sequencing and bioinformatics approach |



- P-043 **Nikolett Virág**
A new cell-autonomous role for the HECA orthologue, Headcase, in blood cell progenitor maintenance in the *Drosophila* lymph gland
- P-044 **Alexandra Gráf**
Comparative analysis of BRCA1 variants of uncertain significance captured by molecular diagnostic testing
- P-045 **Bernadette Hotzi**
Studying the role of an unknown protein in premature ageing in *Caenorhabditis elegans* model organism
- P-046 **Zoltán Kovács**
Differential condensation features of bacterial and eukaryotic nucleoprotein coacervates reflect adaptations to genome metabolic contexts
- P-047 **Regina Martinek**
Assessment of the mutagenic effect of Group 2A pesticides using a whole genome sequencing approach
- P-048 **Ádám Sánta**
Investigation of the effects of small molecules that inhibit PCNA monoubiquitination in human cells
- P-049 **Bendegúz Szabó**
Analysis of genomic rearrangement length distribution in different DNA damage tolerance pathway mutants
- P-050 **Alexandra Fanni Szalmás**
Characterization of protein kinase and phosphatase activity in human pulmonary planocellular and adenocarcinoma
- P-051 **Zoltán Szeltner**
Probing Telomeric-like G4 Structures with Full or Partial 2'-deoxy-5-hydroxyuridine Substitutions
- P-052 **Bernadett Szikriszt**
Testing the effect of human germline BRCA1 mutations on genome stability in a cell line model
- P-053 **Eszter Bajtai**
Characterization of chemotherapy-surviving tumor cells and understanding their role in relapse
- P-054 **Zita Bognár**
Desethylamiodarone may have cytostatic potential on lung adenocarcinoma
- P-055 **Luca Fodor-Varga**
The effects of sodium glucose cotransporter inhibitors on intestinal motility
- P-056 **Balázs Gombos**
Application of transgenic mouse models for chemotherapy optimization