

4th INTERNATIONAL CONFERENCE ON PLANT BIOLOGY 23rd SPPS Meeting







6-8 OCTOBER 2022 BELGRADE

Serbian Plant Physiology Society

Institute for Biological Research "Siniša Stanković" National Institute of Republic of Serbia, University of Belgrade

Faculty of Biology, University of Belgrade

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4th International Conference on Plant Biology (23rd SPPS Meeting)

6-8 October, Belgrade

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THURSDAY 6 [™] OCTOBER		
12:00-18:00	Registration	
12:00-14:00	NEPETOME project workshop (Science Fund of the Republic of Serbia, #Grant No 7749433): "Methodologies for the iridoid diversity investigation within the genus Nepeta" (Botanical Garden "Jevremovac")	
18:00-22:00	Welcoming cocktail and Celebration of SPPS jubilee (Botanical Garden "Jevremovac")	

FRIDAY 7TH OCTOBER

09:00-09:15 *Opening Ceremony*

SECTION 2 · PLANT STRESS PHYSIOLOGY

Chairs: Jelena	Chairs: Jelena Dragišić Maksimović & Tamara Rakić		
09:15-10:00	Keynote: Mondher Bouzayen Uncoupling fruit softening from fruit ripening: a paradigm shift of thinking		
10:00-10:30	Plenary lecture: Miroslav Lisjak Growth conditions may affect the nutritional quality of wheatgrass (Triticum aestivum L.)		
10:30-11:00	Plenary lecture: Hermann Heilmeier The functional role of non-essential elements in the root zone: how interactions between essential and non-essential elements shape the chemical rhizosphere environment		
11:00-11:30	Coffee break		
11:30-11:50	Invited talk: Zsófia Bánfalvi Regulation and function of GIGANTEA genes in Solanum tuberosum cultivar 'Désirée'		
11:50-12:10	<i>Invited talk:</i> Ingeborg Lang Drought or heavy metals – investigating the abiotic stress tolerance in bryophytes		
12:10-12:30	<i>Invited talk</i> : Biljana Kukavica <i>Flooding and antioxidative response in plants</i>		
12:30-12:50	Invited talk: Sonja Milić Komić Distinctive regulation of different phenolics biosynthesis by high light and UV-B in three basil varieties		
12:50-13:05	Selected talk: Mariana Stanišić What happens with phloretin in plants? – Phloretin real-time effects and post-treatment metabolism in treated Arabidopsis seedlings		
13:05-13:20	Selected talk: Danijela Arsenov Fullerenol (C60(OH)24) as a potent stress alleviator against drought and trace-element toxicity in Alliaria petiolata (M.Bieb.) Cavara et Grande		
13:20-14:00	Poster session		
14:00-15:30	Lunch break		
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SECTION 1 · PLANT GROWTH, DEVELOPMENT, METABOLISM AND NUTRITION

Chairs: Ivana Maksimović & Slavica Ninković		
15:30-16:00	Plenary lecture: Panagiotis Kalaitzis A prolyl-4-hydroxylase and Arabinogalactan proteins are involved in relocation of tomato abscission zone	
16:00-16:30	Plenary lecture: Marjorie Guichard State-dependent protein interaction networks of a central regulator of plant growth and metabolism	
16:30-16:50	Invited talk: Václav Motyka Hormonome and role of desiccation in somatic embryogenesis of conifers	
16:50-17:20	Coffee break	
17:20-17:40	Invited talk: Julien Pirrello Transition to ripening in tomato fruit needs genetic reprogramming initiated in gel tissue	
17:40-18:00	Invited talk: Guido Grossmann Robust yet adaptive - morphogenesis and growth regulation in roots	
18:00-18:20	Invited talk: Jan Fíla The beta-subunit of nascent polypeptide associated complex plays a role in flowers and siliques development of Arabidopsis thaliana	
18:20-18:35	Selected talk: Kiril Mishev The interaction network of the plant NudC family protein NMig1	
18:35-19:15	Poster session	
SATURDAY 8 [™] OCTOBER		
09:00-10:00	SPPS Assembly	
	SECTION 4 · ECOLOGY, GENETICS AND EVOLUTION OF PLANTS	
Chairs: Branislav Šiler & Sanja Manitašević Jovanović		

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10:00-10:30	Plenary lecture: Velemir Ninković Plant signaling and behavior mediated via volatiles	
10:30-11:00	Plenary lecture: Janez Kermavnar Impacts of forest management on plant functional traits and ecological conditions in the Dinaric fir-beech forests (Slovenia)	
11:00-11:30	Coffee break	
11:30-11:50	Invited talk: Ksenija Jakovljević Ecophysiology of metal-hyperaccumulation in plants: what do we know so far?	
11:50-12:10	Invited talk: Jelena Milojević Elucidation of the mechanism underlying somatic embryo induction in spinach	

12:10-12:30	Invited talk: Miroslava Zhiponova Catmint (Nepeta nuda L.) Phylogenetics and Metabolic Responses in Variable Growth Conditions
12:30-12:50	Invited talk: Neda Aničić Progress in disentangling the diversity of iridoids within the genus Nepeta: surprising biosynthetic and evolutionary insights
12:50-13:05	Selected talk: Denitsa Teofanova Distribution, host range, and genetic variability of the holoparasitic genus Cuscuta in Bulgaria
13:05-13:20	Selected talk: Katarina Hočevar Variation in Hsp70 and Hsp101 levels in response to experimental warming in Iris pumila L.: an open-topped chamber experiment
13:20-14:00	Poster session
14:00-15:30	Lunch break

SECTION 3 · APPLICATION IN AGRICULTURE, PHARMACY AND FOOD INDUSTRY

Chairs: Ana Ćirić & Ana Marjanović Jeromela		
15:30-16:00	Plenary lecture: Angelos K. Kanellis Aroma formation in Vitis vinifera grape berries	
16:00-16:30	Plenary lecture: Ekaterina-Michaela Tomou Metabolomic strategy for detecting herbal products' differentiations and potential adulteration	
16:30-16:50	Invited talk: Mila Grahovac Essential oils and hydrolates in control of plant pathogens	
16:50-17:20	Coffee break	
17:20-17:40	Invited talk: Carla Vogt Determination of elements, isotopes and organics in plants with high local resolution by mass spectrometric methods	
17:40-18:00	Invited talk: Milan Mirosavljević Integrating physiological traits in local small grains breeding program	
18:00-18:20	Invited talk: Nada Ćujić Nikolić Chokeberry, from natural polyphenol resource to promising functional foods and pharmaceuticals	
18:20-18:35	Selected talk: Ana Pantelić Late embryogenesis abundant (LEA) proteins in Ramonda serbica Panc identification, classification and structural characterization	
18:35-18:50	Selected talk: Dejan Stojković Supercritical fluid extraction of Chicory reveals its antimicrobial, antibiofilm and wound healing potentials	
18:50-19:15	Poster session	
19:15-19:30	Closing Ceremony	
20:00-00:00	Gala Dinner	

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Progress in disentangling the diversity of iridoids within the genus *Nepeta*: surprising biosynthetic and evolutionary insights

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<u>Neda Aničić</u>¹, Dragana Matekalo¹, Marijana Skorić¹, Jelena Božunović¹, Jasmina Nestorović Živković¹, Uroš Gašić¹, Milica Milutinović¹, Slavica Dmitrović¹, Boban Anđelković², Milena Dimitrijević³, Milan Dragićević¹, Danijela Mišić¹

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Nepeta L. is the largest genus of the Lamiaceae family, and the exclusive representative of the subfamily Nepetoidae which produces iridoids. However, the genus Nepeta comprises both taxa producing iridoids (iridoid aglycones-IAs and/or iridoid glucosides-IGs), and those lacking iridoids. Combining omics-guided approach and functional genomics in elucidating the genetic background of the iridoid diversity within the genus Nepeta, we provide novel insights into molecular mechanisms driving the iridoid biosynthesis, and thus into chemical evolution of iridoids within the genus Nepeta. Multidisciplinary experimental approach is here explained on the example of iridoid synthases (IS), functionally characterized by in vitro enzymatic assays using recombinant proteins heterologously expressed in Escherichia coli. Functional ISs are expressed in both iridoid-producing N. rtanjensis (NrIS2) and N. sibirica L. (NsIS), as well as in iridoid-lacking N. nervosa L. (NnIS). N. nervosa possesses an iridoid biosynthetic platform, but inactive one, most likely due to the "switching off" or "silencing" of some gene(s) upstream in the pathway, which suggests that this species has lost the ability to produce iridoids during the evolution. Furthermore, comparative iridoid profiling and co-expression analysis of biosynthetic genes and transcription factors (TFs) during the exposure of chemodiverse Nepeta taxa to environmental stresses (e.g. dehydration, UV-B radiation, pathogens) or elicitors (MeJA), facilitated the explanation of regulatory mechanisms determining the productivity of plants, as well as of factors responsible for the presence/ absence of iridoids or specific groups of iridoids (IAs and IGs), in these plants.

Keywords: Nepeta, dehydration, functional characterization of genes, iridoid synthase, nepetalactone

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