BOOK OF ABSTRACTS



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VRNJAČKA BANJA • SERBIA







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Abstracts of the 6th CONGRESS OF THE SERBIAN GENETIC SOCIETY

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WELCOME TO VI CONGRESS OF THE SERBIAN GENETIC SOCIETY!

Dear colleagues,

Welcome to the 6th Congress of the Serbian Cenetic Society. The Serbian Cenetic Society (SCS) has been founded in 1968 and the first Congress organized by the SCS was held in 1994 in Vrnjacka Banja. Since then, the Congress of Serbian Genetic Society is held every five years. Over the past years, the Congress has grown from a national to an international meeting.

The experience of the past meetings motivated our efforts to continue with this series with a clear tendency to strengthen the scientific connections among researchers from different European countries.

The Congress will focus on the most recent advances in genetics and on wide range of topics organized in 9 sessions and two workshops. Many of the presentations will be in lecture-like settings, but we hope that there will also be ample opportunities for informal interaction outside the scheduled

The successful organization of the Congress has required the talents, dedication and time of many members of the Scientific and Organizing committees and strong support from our sponsors. I hope that you will find the Congress both pleasant and valuable, and also enjoy the cultural and natural beauty of Vrnjacka Banja.

Yours sincerely,

Branka Vasiljevic President of the Serbian Genetic Society

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PLENARY LECTURES

SESSION 1	Human omics variation
SESSION 2	Medical genetics
SESSION 3	Genetic toxicology: from cell to ecosystem
SESSION 4	Adaptation and ecological genetics
SESSION 5	Genetic diversity, phylogeny and conservation
SESSION 6	Breeding for changing environments
SESSION 7	Microbial genetics
SESSION 8	Bioinformatics and big data analysis
SESSION 9	Miscellaneous topics
WORKSHOP 1	Personalized medicine: promise and reality
WORKSHOP 2	The truth is in wine and DNA -applications of molecular methods in viticulture

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06 - 19 Poster

ASSESSMENT OF DURUM WHEAT CULTIVARS BASED ON MORPHOLOGICAL TRAITS

<u>Verica Takač,</u> Sanja Mikić, Milan Mirosavljević, Vojislava Momčilović, Dragana Trkulja, Ankica Kondić-Špika, Ljiljana Brbaklić

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Understanding genetic diversity of durum wheat germplasm is essential for its identification and efficient use in breeding programs. The goal of this study was to assess genetic diversity of durum wheat cultivars from the Institute of Field and Vegetable Crops, Serbia, with 26 morphological characteristics based on International Union for Protection of New Varieties of Plants guidelines. The Shannon diversity index was used as an indicator of morphological diversity and it varies from 0.283, for glaucosity of lower side of the flag leaf blade and density of hairiness of uppermost node of the culm, to 0.950, for the ear color, with the mean value of 0.616, indicating a medium to high level of morphological diversity. On average, the diversity was higher for traits relating to generative organs than for those associated with vegetative organs. The 21 morphological characteristics were sufficient to distinguish unique profiles of all durum wheat varieties. The estimation of varietal diversity and identification of morphological characteristics with the highest discriminative power were done by multiple correspondence analysis. The traits that contributed the most to the distinction of varieties were the ear coloration, length of beak of the lower glume, lower glume shape ear length of awns at tip relative to ear length and color of awns. Morphological characterization using the traits with the highest discriminative power could be a useful complementary method for durum wheat germplasm classification and diversity analysis.

CATEGORICAL DATA, DIVERSITY, DISCRIMINATIVE POWER, MULTIPLE CORRESPONDENCE ANALYSIS, TRITICUM TURGIDUM SUBSP. DURUM

06 - 20 Poster

THE CHEMICAL AND PROTEIN COMPOSITION OF GRAINS OF TWELVE DIFFERENT SWEET MAIZE GENOTYPES (ZEA MAYS L. SACCHARATA)

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Sweet maize grains are used in human nutrition in the endosperm milk stage. The quality of these tender, succulent and sweet grains is determined by genes which distinguish sweet maize from common maize. In recent years, the practice of using sweet maize has been increasing both in our country and worldwide, while its production has therefore been growing. The largest part of sweet maize produced in our country is processed and exported as frozen grains. The most important quality parameters of sweet maize grains are taste, composition of the endosperm, and tenderness of the pericarp.

The paper presents the results of chemical and protein compositions of grains of 12 different sweet maize genotypes in the endosperm milk stage. Since sweet maize grain is used for human consumption, this stage is also a stage of grain harvest maturity. The chemical composition of grains of selected sweet maize genotypes was analysed by determining the content of non-structural carbohydrates (NFC - Non-Fibre Carbohydrate), protein, oil, crude fibre, and NDF (Neutral Detergent Fibre – fibres insoluble in neutral detergent made of hemicellulose, cellulose and lignin). The protein composition of grains of selected sweet maize genotypes was analysed by determining the contents of albumin, globulin, α -zein and glutelin. The dry matter content was established refractometrically and it ranged from 20.1 to 26.0% in observed sweet maize genotypes. The results obtained on the chemical composition of grains of the selected sweet maize genotypes showed that contents of NFC, protein, oil, crude fibre, ash and NDF ranged from 69.64 to 77.29%; 10.26 to 11.98%; 3.81 to 6.89%; 1.36 to 2.28%; 1.73 to 2.39%, and from 6.75 to 9.32%, respectively. The contents of albumin, globulin, α -zein and glutelin ranged from 22.59 to 37.10%, 4.88 to 7.37%, 16.97 to 24.71%, and from 18.18 to 22.55%, respectively.

SWEET MAIZE, CHEMICAL AND PROTEIN COMPOSITION

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