

BOOK OF ABSTRACTS

XIII International Scientific Agriculture Symposium "AGROSYM 2022" October 6-9, 2022



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Jahorina, October 06 - 09, 2022

Impressum

XIII International Scientific Agriculture Symposium "AGROSYM 2022" **Book of Abstracts Published by** University of East Sarajevo, Faculty of Agriculture, Republic of Srpska, Bosnia University of Belgrade, Faculty of Agriculture, Serbia Mediterranean Agronomic Institute of Bari (CIHEAM - IAMB) Italy International Society of Environment and Rural Development, Japan Balkan Environmental Association (B.EN.A), Greece CDR, University of Natural Resources and Life Sciences (BOKU), Austria Perm State Agro-Technological University, Russia Voronezh State Agricultural University named after Peter The Great, Russia Tokyo University of Agriculture, Japan Faculty of Agriculture, University of Western Macedonia, Greece Chapingo Autonomous University, Mexico Selcuk University, Turkey University of Agronomic Sciences and Veterinary Medicine of Bucharest, Romania Slovak University of Agriculture in Nitra, Slovakia National University of Life and Environmental Sciences of Ukraine, Kyiv, Ukraine Saint Petersburg State Forest Technical University, Russia University of Valencia, Spain Tarbiat Modares University, Islamic Republic of Iran Valahia University of Targoviste, Romania Faculty of Bioeconomy Development, Vytautas Magnus University, Lithuania Faculty of Agriculture, University of Akdeniz - Antalya, Turkey Ukrainian Institute for Plant Variety Examination, Kyiv, Ukraine Institute of Animal Science- Kostinbrod, Bulgaria National Scientific Center "Institute of Agriculture of NAAS", Kyiv, Ukraine Department of Agricultural, Food and Environmental Sciences, University of Perugia, Italy Watershed Management Society of Iran Faculty of Agriculture, Cairo University, Egypt Higher Institute of Agronomy, Chott Mariem-Sousse, Tunisia SEASN - South Eastern Advisory Service Network, Croatia Faculty of Economics Brcko, University of East Sarajevo, Bosnia and Herzegovina Biotechnical Faculty, Montenegro Institute of Field and Vegetable Crops, Serbia Institute of Lowland Forestry and Environment, Serbia Institute for Applied Science in Agriculture, Serbia Agricultural Institute of Republic of Srpska - Banja Luka, Bosnia and Herzegovina Maize Research Institute "Zemun Polje", Serbia Faculty of Agriculture, University of Novi Sad, Serbia Institute for Animal Science, Ss. Cyril and Methodius University in Skopje, Macedonia Serbian Academy of Engineering Sciences, Serbia Balkan Scientific Association of Agricultural Economics, Serbia Institute of Agricultural Economics, Serbia **Editor in Chief** Dusan Kovacevic **Tehnical editors** Sinisa Berjan Milan Jugovic Noureddin Driouech Rosanna Ouagliariello Website: http://agrosym.ues.rs.ba

CIP - Каталогизација у публикацији Народна и универзитетска библиотека Републике Српске, Бања Лука

631(048.3)(0.034.4)

INTERNATIONAL Scientific Agricultural Symposium "Agrosym 2022" (13 ; Jahorina) Book of Abstracts [Електронски извор] / XIII International Scientific Agriculture Symposium "Agrosym 2022", Jahorina, October 06 - 09, 2022 ; [editor in chief Dušan Kovačević]. - East Sarajevo =Istočno Sarajevo : Faculty of Agriculture =Poljoprivredni fakultet, 2022. - 1 електронски оптички диск (CD-ROM) : текст, слика ; 12 cm

Системски захтеви: Нису наведени. - Насл. са насл. екрана. - Регистар.

ISBN 978-99976-987-2-8

COBISS.RS-ID 136884481

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BIOINOCULANTS IN SUSTAINABLE AGRICULTURE – THE INFLUENCE ON MAIZE CHARACTERISTICS AND PHYTOBIOME

Ivica DIMKIĆ^{1*}, Aleksandra JELUŠIĆ², Matjaž HLADNIK³, Tamara JANAKIEV¹, Dunja BANDELJ³, Slavoljub VUKIĆEVIĆ⁴, Djordje FIRA¹

¹Chair of Biochemistry and Molecular Biology, University of Belgrade - Faculty of Biology, Belgrade, Serbia
²Department of Life Sciences, University of Belgrade - Institute for Multidisciplinary Research, Belgrade, Serbia
³Department of Applied Natural Sciences, University of Primorska - Faculty of Mathematics, Natural Sciences and Information Technologies (UP FAMNIT), Koper, Slovenia

⁴Fertico D.O.O., Nitrogenous Fertilizer Manufacturing Enterprise, Indjija, Serbia *Corresponding author: ivicad@bio.bg.ac.rs

Abstract

Maize (Zea mays L.) occupies a significant part of the world cultivated area. Beneficial microorganisms and organic fertilizers are powerful tools for sustainable agricultural management. Applying appropriate formulation either through seed coating or composting material together with seeds during the sowing was aimed to achieve optimal performance on plants and increase the crop yields. Also, a complete analysis of phytobiome including highthroughput amplicon and whole-genome sequencing, and computational biology of the treated and non-treated plants' roots and soil was applied. The statistically significant difference in bacterial communities between phenophases and treatments was much more prevalent in the roots than soil itself. In general, seedling phenophase (II) differed in terms of diversity from bacterial communities of flowering (III) and harvesting phenophases (IV), regardless of treatment, indicating good colonization with the treated substrate for up to one month. In the root samples, the dominance of the genus Pseudomonas was evident, in the initial stages (phase II), while this percentage decreased linearly over time. High relative abundance of the genera Achromobacter, Sphingobacterium, and Stenotrophomonas also characterized Π phenophase. Contrary, a relative abundance of the genus Bacillus was characteristic for III phenophase, and especially twice or three times higher in treatments with phytobiotic or manure compared to the negative control. Furthermore, it was observed that with increasing inoculation of Bacillus isolate, the percentage of Pantoea genus decreases drastically in coated seeds. In conclusion, a statistically significant increase in maize yield was obtained in the treatment with phytobiotic with an average value of 650 kg/ha compared to the negative control.

Keywords: Maize, Phytobiotic, Poultry manure, Next Generation Sequencing, Phytobiome.