NOVEL APPROACHES AND TECHNOLOGIES FOR CURRENT AND FUTURE CHALLENGES IN AGRICULTURAL CHEMISTRY

14 – 17 February 2022 – Udine, Italy



Monday, 14 February 2022		
09:00 – 09:10	Opening Session	
Session 1: Introduction to agroecosystem		
Moderators: Gianpiero VIGANI		
09:10 – 09:50	Innovative analytical approaches in soil science: What benefits from x-ray analyses? Concetta Eliana GATTULLO, University of Bari, Italy	
09:50 – 10:30	Introduction to metabolomics and it's potential uses in plant ecophysiology Fabrizio ARANITI, University of Milan, Italy	
Coffee break		
10:45 – 11:45	Structure, function and host control of the crop microbiota <i>Davide BULGARELLI, University of Dundee, Scotland, UK</i>	
	Workshop: Presenting your research	
Moderator: Fabrizio ARANITI		
11:45 – 12:00	Tomato plant responses induced by sparingly available inorganic and organic phosphorus forms are modulated by strigolactones Veronica SANTORO, University of Turin, Italy	
12:00 – 12:15	A collagen-derived protein hydrolysate as plant biostimulant: the importance of elucidating the roles of bioactive peptides Stefano AMBROSINI, University of Verona, Italy	
12:15 – 12:30	Root exudates reuptake and alteration of carbon isotope fractionation by tomato plants under phosphorus deficiency Fabio TREVISAN, Free University of Bozen, Italy	
12:30 – 12:45	Miscible displacement of potassium in the rhizosphere Aline BATISTA, University of São Paulo, Brazil	

Tuesday, 15 February 2022		
Session 2: Plant session		
Moderators: Anita ZAMBONI, Laura ZANIN		
09:00 – 10:00	Machine learning to get insights into nitrogen signaling interactions in plants Gabriel KROUK, SupAgro Montpellier, France	
10:00 - 10:40	Effect of microbial communities on gene expression in plants <i>Fabio MARRONI, University of Udine, Italy</i>	
Coffee break		
11:00 – 11:40	The potential of MS-based untargeted metabolomics in plant science Maria Begona MIRAS MORENO, University of Piacenza, Italy	
11:40 – 12:20	Photomodulation in plant biology: applications, perspectives and opportunities Marco LANDI, University of Pisa, Italy	
12:20 – 12:30	Sponsor Time	

Wednesday, 16 February 2022		
Session 3: Soil session		
Moderators: Daniela PEZZOLLA, Daniel SAID PULLICINO		
09:00 - 10:00	Plant- or microbial-derived? Sources of soil organic matter and their relevance to carbon dynamics Gerrit ANGST, Biology Centre CAS, Czech Republic	
10:00 - 10:40	Developing novel Lunar and Martian regolith-based substrates for pedogenesis and space farming studies Antonio CAPORALE, University of Naples, Italy	
Coffee break		
11:00 – 11:40	Microsensor technology to quantify acclimation traits in rice roots against soil anoxia Elisa PELLEGRINI, University of Udine, Italy	
11:40 – 12:20	X-ray Computed Tomography for Soil and Plant Sciences Applications Craig STURROCK, University of Nottingham, UK	
12:20 – 12:30	Sponsor Time	

Thursday, 17 February 2022		
Session 4: Microorganism session		
	Moderators: Youry PII, Nicola TOMASI	
09:00 – 09:40	Omics approaches to study soil functionality Laura GIAGNONI, University of Brescia, Italy	
09:40 – 10:20	The hidden role of soil biodiversity in plant and animal health: Making the invisible visible Luigimaria BORRUSO, Free University of Bozen, Italy	
Coffee break		
Workshop: Presenting your research		
	Moderator: Sara BUOSO, Antonio CAPORALE	
10:40 – 11:40	From data to stories: public science communication is a complex and fascinating challenge <i>Elisabetta TOLA, Formicablu, Italy</i>	
11:40 – 11:55	Microclimate, soil chemistry and microbiota fail to explain Janzen-Connell distribution of <i>Euphorbia dendroides</i> in a Mediterranean shrubland Mohamed IDBELLA, University of Naples Federico II, Italy	
11:55 – 12:10	Impact of tillage erosion state on ¹⁴ C labelled assimilate C distribution in a spring rapeseed (<i>Brassica napus</i> L.) soil system and soil C pools Ayten PEHLIVAN, Leibniz-Centre for Agricultural Landscape Research and Humboldt University of Berlin, Germany	
12:10 – 12:25	Assessing phosphorus availability and uptake by paddy rice: soil testing and plant responses Sara MARTINENGO, University of Turin, Italy	
12:25 – 12:40	An integrative approach to study the responses of <i>Zea mays</i> L. to organic nitrogen provided to roots in the form of amino acids <i>Chiara MURATORE, University of Milan, Italy</i>	
12:40 – 13:00	Best Presentation Awards & Closing session	

An integrative approach to study the responses of *Zea mays* L. to organic nitrogen provided to roots in the form of amino acids.

Muratore C., Araniti F., Espen L., Prinsi B.

Department of Agricultural and Environmental Sciences - Production, Landscape, Agroenergy, University of Milan, via Celoria 2, 20133 Milano, Italy

Keywords: Plant nitrogen nutrition, Amino acids, Proteomics.

Although the capacity to take up organic nitrogen (N) in the form of amino acids is widespread among plants, their relevance for N nutrition in crops has been little investigated compared to inorganic N sources. In the last years, several studies have pointed out that distinct amino acids exert different effects on plant morphology and physiology. However, less is known about their effects on plant metabolism, especially when supplied to the roots at composition and concentration mimicking soil conditions. In this respect, this work aimed to investigate the capacities of crops, such as maize (Zea mays L.), to utilize amino acids as N nutrients through an approach that integrates morphological, physiological, biochemical and proteomic evaluations. Maize seedlings were grown by a hydroponic system for three days in the absence of N and then were provided for further four days with 250 µM of a mixture of eight amino acids [Glutamic acid (50 μM), Glutamine (50 μM), Glycine (50 μM), Alanine (50 μM), Aspartic acid (12.5 μM), Arginine (12.5 μ M), Lysine (12.5 μ M) and Threonine (12.5 μ M)], or with 250 μ M of inorganic N in the form of nitrate. Plants were analyzed after five and seven days of hydroponic growth to follow both early responses and later metabolic adjustments. Plants supplied with amino acids continued to grow over the experiment, although less than those supplied with nitrate. Therefore, plants utilized both the N sources for growth. In detail, amino-acid treated plants were characterized by a decreased root/shoot ratio and a shorter but thicker root system than nitrate-treated plants. The responses at a nutritional and metabolic level significantly differed, as suggested by changes in the plant content of nitrate, ammonium and key metabolites such as reducing sugars and sucrose. The LC-ESI-MS analysis of amino acid composition in roots, xylem sap and leaves revealed that the roots of plants supplied with amino acids showed an upsurge in the levels of uncharged and basic amino acids. Interestingly, glutamine and alanine were the major amino acids translocated through xylem sap for both the nutritional conditions, while the proportion of amino acids in leaves was more constant over time. Overall, these results highlight the different roles played by roots and leaves in response to N inputs and suggest that roots perceived and adapted promptly to the kind of N available and influenced N translocation to the shoot. In order to clarify the meaning of these metabolic adjustments, large-scale analysis of the total proteome fraction of roots and leaves based on onedimensional Gel Liquid Chromatography-Mass Spectrometry (1D GeLC-MS/MS) is ongoing. This study will improve the knowledge about plant responses to different N inputs and plant metabolic use of amino acids, providing information that could be useful for enhancing the sustainability of crop production.