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## CONFERENCE PROGRAM AND ABSTRACTS



Dynamics of Biogeochemical Systems: Processes and Modeling

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## Metabolic and genetic profile of selected heat-resistant Byssochlamys strains and its decimal reduction time (D value) at 80°C and 95°C

K. Oszust<sup>(1)</sup>, M. Frąc<sup>(1)</sup>, N. Bilińska-Wielgus<sup>(1)</sup>, A. Gryta<sup>(1)</sup>, A. Pawlik<sup>(2)</sup>, G. Janusz<sup>(2)</sup> and M. Piotrowska<sup>(3)</sup>

(1) Institute of Agrophysics of the Polish Academy of Sciences, Lublin, Poland
(2) Maria Curie-Skłodówska University, Department of Biochemistry, Lublin, Poland
(3) Institute of Fermentation Technology & Microbiology, Lodz University of Technology, Lodz, Poland

Species of the genus *Byssochlamys* occur in different environment, including soil and cause the spoilage of agricultural raw materials, fruit and foods due to contact with soil. Some of them produce heat-resistant ascospores and synthetize toxic patulin. Therefore they are dangerous for human health. On the one hand the knowledge of metabolic and genetic profile of fungi can be useful in determination of active compounds for new fungicides [1]. On the other hand metabolic pattern can be useful in the evaluation of fungal potential in biogeochemical application such as participation prediction and assessment in transformations processes occurring in soil environment.

The aim of the study was the evaluation of metabolic pattern and genetic profile of 3 *B. fulva* and 3 *B. nivea* strains. The study included also the assessment of decimal reduction time (D-value) of strains at 80°C and 95°C. Temperatures were chosen as the most frequently used in fruit products processing in pasteurization.

Metabolic profile was evaluated using FF plates (Biolog<sup>TM</sup>). We obtained data on utilization of 95 carbon sources from different groups: carbohydrates, amino acids, amines and amides, polymers, carboxylic acids and miscellaneous substrates. All *Byssochlamys* strains were genetically characterized using the amplified fragment length polymorphism (AFLP) technique according to Vos et al. [2] with modification made by Pawlik et al. [2]. D values for each strain were evaluated using plate methods with 10 days incubation at 28°C.

The Biolog™ FF analysis allowed comparison of functional diversity of the *B. fulva* and *B. nivea* fungal strains. Significant differences have been demonstrated in substrate richness values. *Byssochlamys* strains included in the analysis displayed an AFLP profile in various similarity level. Higher differences in D-values between studied strains were observed during incubation at 80°C. D-values for all studied strains at 95°C were below 1 minute.

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<sup>[1]</sup> M. Frac, S. Jezierska-Tys, T. Yaguchi, Advances in Agronomy 132, 2015, 161-204.

 <sup>[2]</sup> P. Vos, R. Hogers, M. Bleeker et al., Nucleic Acids Research, 23(21), 1995, 4407-4414.
 [3] A. Pawlik, G. Janusz, I. Dwbska, M. Siwulski, M. Frac, J. Rogalski, BioMed Research International, 2015, doi.org/10.1155/2015/726149