28 - 31 October 2020

ELECTRONIC ABSTRACT BOOK



in association with the Serbian Society of Microbiology





28 - 31 October 2020



Message from the organizers

Dear solleagues and friends,

We would like to confailly great you at the end of the PEMS 2020 ONLINE Conference on Microbiology, that has been successfully organized at the end of October 2020.

One great idea became reality it would not have been possible without all of you, for which we are grateful. A planeering endeavor in many different aspects and in challenging times. First, as a first Conference of the kind in the FEMS history, planned as a middle sized event, oriented in parallel to regional and global audiences, both in terms of content delivery and participation and an event done in partnership with the national/regional association and the dedicated PCD. Second, as a first major Conference event fully organized as an ONUNE format in European microbiology in 2020. Third, as the first major FEMS Conference event organized in complex times, with numerous challenging, even adverse psychological, societal and economic effects for the whole professional community, and, in addition, inscribiologists being the one of the few medical professions directly standing in the front line of the COVID-19 response

We made a brave decision and it paid off. With the new world of ONUNE possibilities, we strive and succeed to organize the most suitable formats to cover the scientific content of the conference, discuss the most challenging topics of the day and to enable scientific exchange and the interaction with the participants in a select possible manner

We are happy to say that, despite all different complexities, overall results of the Conference and the whole concept went beyond our bravest expectation. Almost 700 recrobiologists from 40 European countries and another 20 worldwide, making the ONUNE Conference global event, eight scientific sessions, including one session with best selected grantees presentations and four Symposiums, organized in two parallel streams, four plenary lectures and COVID-19 Round Table for the whole audience, total of over 70 insited lectures with prereconded lectures. In addition, over 50 prereconded presentation prepared by participants-authors in the main program, in 7 corresponding ORAL CMLINE sessions, all together more than 30 hours of prerecorded material. Finally, another 60 video presentations, over 400 e-posters and some 10 Company materials, in total almost 600 presentational items, uploaded on the Conference DMUNE platform and accessible to participants until the 27. October 2021.

As FEMS, we are delighted that we were able to put together a great scientific program and contribute to the Conference success. Moreover, in Spht of the challenging times we live in and all different complexities it entails, and also to support your enthusiasm and scientific rigous; we decided to support your effort by waiving conference fees for the ONLINE Conference for almost 300 collegaues, authors of scientific papers, submitted until 15, March 2020.

Despite the circumstances, the pharmaceutical and medical indicatry has been with the Conference, providing valuable

We sincerely hope that you also enjoyed all different contents and aspects of the Conference and that you would be able to catch up with those recordings and presentations you may have missed or have a detailed look at posters

We would like to warmly with you health, love and happiness, Merry Christmas and a Happy New Year and are looking forward to the new encounters, in the next occasion, FEMS 2022 Conference in Belgrade.

Prof. dr Hilary Lappin Scott

Sincerely

Maseur.

Prof. & Vaso Taleski, Chair, COC

Prof. dr Lazar Ranin, Co-chaig COC

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Prof. dr Dragojio Obradović, Conchete, CSC

Druet

Prof. dr Bauke Oudegs, Charl. CDC



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in association with the Serbian Society of Microbiology

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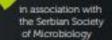
Russia



28 - 31 October 2020

MAIN CONFERENCE PROGRAM





28 - 31 October 2020

WEDNESDAY, 28th October 2020

Schedule		Schedule	
14.00-16.15 Symposium 1 – The Control of Viral diseases by vaccination – new challenges Chairs – Tanja Jovanović, Maja Stanojević		14.00-15.30	Symposium 2 – Helicobacter pylori – microbiologist prospective Chairs – Nataša Opavski
14.00-14.15	Tanja Jovanović, Serbia Viral vaccines – from Jenner to new technologies	1400-1415	Biljana Selimović, Serbia Immunopathogenesis and resistance in H. pylori infection
14.15-14.30	Maja Stanojević, Serbia Poliovirus eradication: novel attenuated poliovirus strains	14.15-14.30	Vladimir Milivojević, Serbia Advances in Helicobacter pylori infec- tion diagnostics and therapy from the gastroenterology comer
14.30-14.45	Maja Čupić, Serbia Traditional and New Influenza Vaccines - the impact of repeated influenca vaccine effectiveness on global health	1430-1445	
14.45-15.00	Ivana Lazarević, Serbia Advances and Challenges in HCV Vaccine Development	14.45-15.00	Mirjana Rajilić-Stojanović, Serbia Gastric microbiota beyond Helicobacter pylori
15.00-15.15	Ana Banko, Serbia MMR vaccine: updates and challenges		1 2 2
15.15-15.30	Aleksandra Knežević, Serbia Future prospects of prophylactic HPV vaccine	15.00-15.30	Discussion and
15:30-15:45	Danijela Miljanović, Serbia Is now the time to introduce VZV vaccine in the mandatory immunization programme in Serbia?		Q6A

15.45-16.15	Discussion and Q&A 15.30-15.40		Break		
16.15-16.25 16.25-18.30	Symposium 3 – Mighty microbes for a healthy and sustainable future Chairs – Tanja Berić, Milan Kojić	15.40-18.20	Symposium 4 – HIVAS – Contemporary approach to the AIDS infection – role of Microbiologist-Virologist Chairs – Jovan Ranin, Danijela Simić		
16.25-16.40	Jasmina Nikodinović Runić, Serbia Plastic degrading microorganisms: Opening an avenue to a sustainable world	15.40-15.55	Danijela Simić, Serbia Epidemiological situation of HIV infection in Serbia and the World		
16.40-16.55	Svetlana Soković, Serbia GABA producing lactobacilli: a way to modulate immune response via autophagy induction	15.55-16.10	Ivana Pesić, Serbia Application of PCR tests in the diagnosis of HIV infection in Serbia: case reports		
16.55-17.10	Mina Obradović, Serbia Bacteriophages - alternative agents for treatment of MDR pathogens	16.10-16.25	Jovan Ranin, Serbia Microbiological principles in contemporary antiretroviral treatment of HIV patients in Serbia		
17.10-17.25	Sanja Jeremić, Serbia Bacterial nanocellulose for biomedical applications: Recent advances and future prospects		Ljiljana Popović, Serbia Dark side of the antiretroviral treatment		
17.25-17.40	Andrea Radalj & Jakov Nišavić, Serbia Some viral infections of animals as potential public health risk	16.40-16.55	Mila Paunić, Serbia An example of good practice - VCCT HIV counseling at the Institute for Students' Health of Belgrade University		
17.40-17.55	Sladana Prišić, USA "Known unknowns" – secret life of Mycobacterium tuberculosis.	16.55-17.15	GlaxoSmithKline Jovan Ranin, Serbia Dolutegravir as core agent in dual therapy		
17.55-18.30	Discussion and Q&A	17.15-17.35	Maja Stanojević, Serbia The Importance of Dolutegravir high barrier to resistance from virological perspective		
		17.35-18.30	Discussion and Q&A		
18.30-19.00	FEMS Networking Session				
19.00	OPENING CEREMONY Chairs: Bauke Oudega, Dragojlo Obradović, Vaso Taleski 19.20 Welcome address – Hilary Lappin-Scott, UK 19.30 Plenary lecture 1 - Anita Kriško, Germany & Miroslav Radman, Croatia/France Survival Strategies in Microbes: it is all about communication 20.30 Plenary lecture 2 - Rino Rappuoli/Mariagrazia, Italy The emerging problem of AntiMicrobial Resistance and the role of vaccines				

	THURSDAY, 29th October 2020				
Schedule					
13.00-14.00	FEMS Networking Session				
14.00-14.40	Chairs – Jozef Anné, Dragana Vuković Tom Coenye, Belgium]			
14.40-14.50	Metabolic adaptations as drivers for ant	Discussion			
14.50-18.10	Session 1 Infections and control (human medical focus) Chairs – Bauke Oudega, Branislava Kocić	14.50-18.10	Session 2 Food Microbiology and Biotechnology (non human medical focus) Chairs – Andriy Sibirny, Miroslav Vrvić		
14.50-15.15	Eliora Ron, Israel Combatting antibiotic resistant septi- cemic <i>E. coli</i>	14.50-15.15	Lixin Zhang, China Harnessing cellular triacylglycerol pool for titer improvement of polyketides		
15.15-15.40	Lidija Šenerović, Serbia Antivirulence agents in the treatment of mixed species infections.	15.15 -15.40	Daniele Daffonchio, Saudi Arabia Microorganisms of the agro-ecosystem, "plant wellness" and the quality of food products.		

_	Ivana Gobin, Croatia Opportunistic premise plumbing pathogens: a new challenge in the hospital Ina Gajić, Serbia Molecular insight into the antimicrobial resistance of group B streptococci, serotype distribution and emerging serotype III sequence type 17	16.05-16.30	Ken Ichi Yoshida, Japan Application of the Gram-positive conjugative plasmid pLS20 for rapid and easy transformation of intra- and inter-species Hristo Najdenski, Bulgaria The prevalence of Yersinia in Europe I would recommend the topic Infections and control Aleksa Obradović, Serbia Novel approaches to biocontrol of plant bacterial diseases.		
16.30-16.55	Danai Pervanidou, Greece West Nile virus in humans in Greece, 2010-2020: surveillance and public health response	16.55-17.20	Andriy Sibirny, Ukraine Non-conventional thermotolerant yeast Ogataea polymorpha for production of the 2nd generation fuel ethanol from lignocellulosics and crude glycerol		
16.55-18.10	Discussion and Q&A	17.20-18.10	Discussion and Q&A		
18.10-18.20	Break	18.10-18.20	Break		
18.40-18.50	Metka Paragi, Slovenia National Laboratory Surveillance of invasive Pneumococcal Diseases (IPD) in Slovenia Discussion and Q&A				
18 50-19 3	FEMS SESSION Patrik Bavoil, France/USA FEMS Director of Publications and Science The FEMS Journals, the new Open Access Journals, and how to publish Patrik Michel Bavoil What makes a great paper? Max Haggblom FEMS Microbes: Open science toward a sustainable world Kimberly Klein FEMS Session Discussion Patrik Michel Bavoil				
19.30-19.40	Break	19.30-19.40	Break		

19.40-21.10	ONLINE Oral Session No. 1	19.40-20.30	ONLINE Oral Session No. 2
19.40-19.45	Diego Emilio Pérez-Stuardo, Chile Lysosomal reactivation in response to IgM-beads on Atlantic salmon macrophages	19.40-19.45	Anwar Kandari, UK Investigation into the effects of fibres and starches on the gut microbiota
19.45-19.51	Gülcan Kuyucuklu, Turkey Effect of N-Acetylcystein-Ampicillin combination against Methicillin resistant Staphylococcus Aureus and their Mec-A gene expressions	19.45-19.50 19.50-19.55	Büşra Akdeniz Oktay, Turkey Potential use of hazelnut husk for pullulan production by Aureobasidium pullulans AZ-6 Dina Ramić, Slovenia New approach for quantification of signal molecules AI-2 of Campylobacter jejuni
19.51-19.59	Lena Ciric, UK Collecting and using evidence for improved hospital cleaning	19 55 -20 00	Izlem Haktanir. France Mechanism of antimicrobial activity of honeybee (Apis mellifera) venom on Escherichia coli and Pseudomonas spp.
19.59-20.04	Miriam Latorre-Millán, Spain Presence of chronic diseases in elderly patients, hospitalized with flu or serious acute respiratory infection	20.00-20.06	Sarah Marie Stoppel, Norway Pre-Treatments and Long-Range PCR to Assess Infectivity of Norovirus Surrogates
20.04-20.16	Mümtaz Güran, Cyprus A survey of oropharyngeal HPV infections among healthy university students: infection prevalence, associated risk factors and awareness levels	20.06 -20.11	Tolulope Modupeoluwa Adeleye, Nigeria Ethanol Production from Cassava Starch by Protoplast Fusants of Wickerhamomyces anomalus and Galactomyces candidum Viorica Corbu, Romania
20.16-20.21	Nataša Stanković Nedeljković, Serbia ACINETOBACTER BAUMANII COMPLEX IN CLINICAL SAMPLES AND ANTIBIOTIC RESISTANCE-DECADES BACK	Screening of newly is showing no pathoge	Screening of newly isolated yeast strains showing no pathogenic profile for biotechnological applications
20.21-20.27	Rare Clinical Presantation caused by <i>Borrelia burgdorferi</i>		
20.27-20.46	5 Yvonne Cortese, Ireland A Novel In Vitro Urethra Model to Visualise Bacterial Migration During Long-Term Urinary Catheter Insertion	20.16-20.3	Discussion and Q&A
20.46-21.1	Discussion and Q&A		

	FRIDAY, 30th	Octobe	r 2020
Schedule	***************************************		
14.00-14.40	Plenary Chair – Vaso Taleski, Dejan Baskić, La Petra Dersch, Germany Remodelling Yersinia-host interactions		rworse?
14.40-14.50	1.023	Discussion	
14.50-17.40	Session 3 AMR, noval therapeutic agents Chairs – Vaso Taleski, Dejan Baskić, Lazar Ranin	14.50-17.40	Session 4 Microbiomes and human health (mixed focus) Chairs – Bauke Oudega, Jelena Begović
14.50-5.15	Stefano Donadio, Italy Scientific and financial challanges for new antibiotics	14.50-15.15	Jeroen Raes Belgium Quantitative microbiome profiling in health and disease
15.15-15.40	Branislava Kocić, Serbia Susceptibility pattern of Gram-negative bacilli to antimicrobial agents with an emphasis on colistin and carbapenems		Özgür Kurt, Turkey Neglected residents of the gut microbiota Nataša Golić, Serbia
15.40-16.05	Rossitza Vatcheva-Dobrevska, Bulgaria Approaches to prevent and control the spread of mdr pathogens in healthcare setting		Gut microbiota and autoimmune diseases Maja Rupnik, Slovenia Clostridium difficile and One Health (position possibility 2)
16.05-16.50	Discussion and Q&A	16.30-16.55	Marina Dinić, Serbia Plasmid-mediated Cephalosporinase in Enterobacterales - Challenges in Detection

16.50-17.00	Break	16.55-17.40	Discussion and Q&A			
17.00-17.50	Lucia Rubio Fernandez, Belgium The European Research Council – funding opportunities for bright minds	17.40-17.50	Break			
17.50-18.30	PFIZER					
	Matteo Bassetti, Italy Novel antibiotics in the era of AMR					
18.30-18.40	Break					
18.40-18.50	PROMEDIA					
	TBD CerTest – Specialists identifying causal pathogens of infectious diseases					
18.50-19.00	Discussion and Q&A					
19.00-19.40	FEMS Networking Session					
19.40-19.50	Break					

19.50-21.20	ONLINE Oral Session No. 3		ONLINE Oral Session No. 4
19.50-19.55	Manuela Agudelo-Restrepo, Colombia Mosquito-larvicidal activity of bacterial extracts produced by Colombian strains	19.50-19.57	Isabel Abellan-Schneyder, Germany Full-length versus short amplicon 16S rRNA sequencing: benefits, drawbacks and risks
19.55-20.00	Tahrioui Ali, France Pistacia lentiscus L. Fruit Thwart Pseudomonas aeruginosa Virulence Through Membrane Fluidity Alteration	19.57-20.06	Jonathan Josephs-Spaulding, Germany PhenoScreen: Development of a Systems Biology Tool for Decision Making in Recurrent UTIs Maria-Isabel Rojas, USA
20.00-20.05	Olukemi Aromolaran, Nigeria Antimicrobial and synergistic effect of green synthesized silver nanoparticles against multi-drug resistant bacteria		Phage as drivers of microbiome evolution: how they regulate their lysogenic-to-lytic switch to move across systems
20.05-20.10	Onyedikachi Cecil Azuama, Eure, France Inhibition of Pseudomonas aeruginosa virulence by Azorella atacamensis from Taira Atacama's community	20.11-20.22	Tomoka Urushidani, Japan Promoting and memory effects of air-solid biofilm culture on persister formation: analysis on mutants of toxin-antitoxin system genes
T-177-178-178-178-178-178-178-178-178-178		20.22-20.57	ONLINE Oral Session No. 5
20.10-20.15	Celia Campillo, Spain Tn-Seq reveals a panel of genes as key contributors to Haemophilus influenzae respiratory infection in a murine model of emphysematous lung	20.22-20.27	IEsra BARAN, Turkey Evaluation of Sysmex UF-5000 for cerebro- spinal fluid and body fluids
20.15-20.27	Nitin Chandra Teja Dadi, Slovakia Antimicrobial properties of a new hybrid system based on clay mineral saponite and photosensitizer phloxine B	20.27-20.32	Nahikari López López, Spain A genome-scale metabolic reconstruction of Haemophilus influenzae reveals FabH as a potential target for antimicrobial thera- peutic development
	Eduardo Lanzagorta García, Ireland Sustainable Antimicrobial Biopolymeric Food & Biomedical Film Engineering Using Bioactive AMP-Ag+ Formulations	20.32-20.37	Özgür Yanılmaz, Turkey Evaluation of fluorescence flow-cytometry to rule out negative urine samples in neonates
	Allison Lopatkin, USA Predicting and inhibiting the spread of plasmid-mediated antibiotic resistance	20.37-20.43	Sayuri Kasagaki, Japan Involvement of DNA methylation in memory phenomenon of persisters
4	Valérian Lupo, Belgium Mining the hidden diversity of class-D (OXA) beta-lactamases	20.43-20.50	in Escherichia coli Ceylan Polat, Turkey
20.54-21.00	Daria Olkiewicz, Poland		Dobrava hantavirus variants in Apodemus flavicollis mice in European part of Turkey
	Antibacterial activity of trans-cinnamaldehyde and geraniol	20.50-20.57	Marko Janković, Serbia Cytomegalovirus glycoprotein B and N genotypes in pediatric recipients of hema- topoietic stem cell transplant
21.00-21.20	Discussion and Q&A	20.57-21.20	Discussion and Q&A
4			

SATURDAY, 31th October 2020

Schedule			
09.00-10.00	FEMS Networking Session		
10.00-12.30	Session 5 Pathogenicity, virulence, diagnosis Chairs – Eliora Ron, Vera Mijač	10.00-11.30	Session 6 Biofilms Chairs – Colin Harwood, Helena Bujdakova
10.00-10.25	Maximiliano Gutierrez, UK Host cell environments and antibiotic efficacy in tuberculosis	10.00-10.25	Ute Römling, Sweden Characterisation of human commensal Escherichia coli strains, Commensal E. coli strains - Unravelling secrets of biofilm formation and beyond
10.25-10.50	Irena Arandelović, Serbia Prediction of susceptibility to first line tuberculosis drugs by DNA sequencing	10.25-10.50	Milan Kojić, Serbia Impact of chromosomal architecture on the response to stress conditions in bacteria
10.50-11.15	Vladimir Peličić, UK Type 4 pill in Neisseria, Streptococcus	10.50-11.15	Helena Bujdakova, Slovakia Photodynamic inactivation - a promising approach combating resistant microorganisms
11.15-11.40	Ivana Čolović-Čalovski & Aleksandar Džamić, Serbia Diagnosis of fungal infection	11.15-11.30	Discussion and Q&A ONLINE Oral Session No. 6

11 40-12 05		roatia ethods in detection and their toxins	11 30-11 38	Gülcan Kuyucuklu, Turkey EVALUATION OF BIOFILM FORMATION IN STAPHYLOCOCCUS STRAINS WITH CONVENTIONAL AND MOLECULAR METHODS
			11.38-11.43	Iago Grobas, UK Swarming Bacillus subtilis transits into a biofilm templated by active jamming in a gradient of antibiotics
			11.43-11.48	Nezahat Koşar, Turkey Acinetobacter baumannii's biofilm formation as a survival and adaptation mechanism
12.05-12.30	Disc	ussion and Q&A	11.48-12.00	Discussion and Q&A
12.30-12.40		Break	12.00-12.40	Break
12.40-13.20	BIOMERIE	UX		
	Valeria Zucch	elli, Italy		
	The role of dia	gnostics in navigating th	rough the challe	nge of COVID-19 pandemic and AMR
13.20-13.30			Break	
	13.30-16.30 COVID-19 ROUND TABLE Moderator – Linda Kenney, USA			
	13.30-13.40	Tanja Jovanović, Serb NEW CORONAVIRUS		EALITY
	13.40-13.50	Miroslav Radman, Cro Viral Parabiosis Hypol		of COVID-19
	13.50-14.00	Ivan Dikić, Germany Ubiquitin networks in	regulation of (COVID-19
	14.00-14.10	Peter Hotez, USA Recombinant Protein	s in COVID19 V	accine Development
	14.10-14.20	Jean-Claude Piffaretti, Switzerland The COVID-19 pandemic seen by the past chair of the Swiss committee for preparing pandemics		
	14.20-14.50	Stefan Pöhlmann, Germany SARS-CoV-2 entry into cells and its inhibition		
	14.30-14.40	Alexander Suvorov, Ro Development of muci		ainst COVID-19
	14.40-14.50	Marion Koopmans, Th Insights in Virus Trans		Genomic Epidemiology
14.50-16.20		Rout	nd Table discu	ssion
16.20-16.30			Break	

16.30-19.30 Session 7 16.30-17.50	Session 8
Environmental Microbiology Chairs – Hilary Lappin-Scott, Branka Vasiljević	FEMS session and short presentations by grantees Chairs - Vaso Taleski, Jozef Anne
16.30-16.55 Cagri Ergin, Turkey World-wide Cryptococcal pathogens	FEMS Grants: Influence on the scientific careers of Early Career Researchers
with respect to global climate change and ecological niche modelling	Rita Hirmondó, Hungary Investigation of the mutator effects of currently used TB drugs provides
16.55-17.20 Vaso Taleski, North Macedonia Metagenomic insight into the microbial diversity of the world unique loran-	insight into the mechanism of drug resistance development in Mycobacteria
dite mine Allchar	Katarina Novović, Serbia Unraveling regulon of novel two-component system in Pseudomonas
17.20-17.45 Colin Harwood, UK The Bacillus secretion factory 17.00-17.10	putida WCS358 through transcriptomics Zandra Fagernäs, Germany
17.45-18.10 Dragan Radnović, Serbia	Biogeography of the ancient oral cavity Kusum Dhakar, Israel
Bacterial resistance to hexavalent chromium and applicability 17.10-17.20 in bioremediation	Modelling of microbial communities to promote atrazine degradation in crop soils
17.20-17.30 18.10-18.35 Ines Sviličić Petrić, Croatia	Amanda Pacholak Poland Azole fungicides – do they affect the properties of environmental bacterial
Microorganism - early indicators of anthropogenic perturbations within the environment	strains?
17.30-17.40	Marija Lješević, Serbia Bacterial degradation of monocyclic aromatic compounds in kraft lignin

	17.40-17.50	Discussion and Q&A
	17.50-18.00	Break
	18.00 -19.30	ONLINE Oral Session No. 7
18.35-19.30	18.00-18.08	Andrea Silva Claros, Spain Heavy metal biosorption using Halomonas strains isolated from the Bolivian altiplano
Discussion	18.08 -18.13	Dragana Tamindžija, Serbia Biodegradation and biosorption of organo- phosphorus pesticides by the alluvial sedi- ment isolate Bacillus megaterium BD5
and Q&A	18.13-18.21	Francesca Bandini, Italy Fate and microbial ecology of biodegradable polymers under industrial anaerobic and aerobic conditions
	18.21-18.28	Guillermo Budiño, Spain Shewanella T2:3D-1.1, an electrifying tale in the deep subsurface
	18.28-18.33	Jose Francisco Nava Olivas, France STUDY OF NAPHTALENE AND PHENANHTRENE DEGRADING BACTERIA ISOLATED FROM SLECTIVE MEDIA
	18.33-18.38	Kalonji Abondance Tshisekedi, South Africa Comparative genomic analysis hints at psychrophilic nature of the novel yeast species Naganishia randhawae

	18.38-18.43	Kusum Dhakar, Israel
		Modelling of microbial communities to promote atrazine degradation in crop soils
18.35-19.30	18.43-18.48	Marija Mojicević, Ireland
		Plastic Saga - The microbes strike back
	18.48-18.53	Sara Fareed Mohamed Wahdan, Ger-
Discussion		many
Discussion		Link between metabolically active soil
and		microbiome and soil function and their response to environmental factors in
and		grassland ecosystem
Q&A	18.53-18.58	Harold Alejandro Jurado Rivadeneira,
1000		Hungary
		Assessing the degradation of short chain chlorinated aliphatic hydrocarbon contaminated sites by monitoring the effect of two different bioremediation approaches using area specific bioaugmentation culture
	18.58-19.06	Karla Peranić, Croatia
		DYNAMICS OF Cryphonectria parasitica
		POPULATIONS AND ASSOCIATED Cry-
		phonectria hypovirus 1 PREVALENCE IN NORTH-WESTERN CROATIA
	19.06-19.11	Munavvara Dzhuraeva, Tajikistan
		THERMOPHILIC BACILLI ISOLATED
		FROM KHOJA OBI GARM (TAJIKISTAN) GEOTHERMAL SPRING
	19.11-19.30	Discussion
	Break	
19.40-19.50		
19.40-19.50		
20.00 CLOSING C		AND ADDRESS Taleski, Lazar Ranin



28 - 31 October 2020

CONFERENCE ABSTRACTS



in association with the Serbian Society of Microbiology

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SESSION 7 - ENVIRONMENTAL MICROBIOLOGY

METAGENOMIC INSIGHT INTO THE MICROBIAL DIVERSITY OF THE WORLD UNIQUE LORANDITE MINE ALLCHAR

Vaso Taleski¹, Ivica Dimkić², Blazo Boev¹, Ivan Boev¹, Sanja Živković³, Slaviša Stanković²

Table 1 and 1 and

- 1 University "Goce Delchev" Shtip, North Macedonia
- 2 University of Belgrade, Faculty of Biology, Serbia
- 3 University of Belgrade, Institute of Nuclear Sciences Vinca, Serbia

Abstract

Background

The Allchar mine situated in southern part of the Republic of North Macedonia, is an unique mine well known as the source of the lorandite (TlAsS2) that present mineral with purest contain of thallium (TI) in the world. The Crven Dol mine, part of the Allchar mine complex, holds an estimated 500 t of Tl. As a result of interaction of solar neutrinos with 205Tl of the lorandite, Tl transforms to 205 Pb (205 lead whose 15,3 million years half-life present traps of neutrino capture event). Lorandite from this ore deposit is used as geochemical detector to capture neutrinos from the Sun, developed in the project LOREX).

Objectives and Methods of this study was, for the first time for this locality, to evaluate presence and diversity of bacteria and fungi using metagenomic methods and bioinformatic analysis (Krona viewer).

Results showed high bacterial and fungal diversity.

Most dominant bacterial phyla level were: Proteobacteria, Bacteroidetes, Acidobacteria, Planctomycetes, Actinobacteria, and Verrucomicrobia. On the genus level following groups of bacteria were detected: Chryseolinea, Opitutus, Flavobacterium, Pseudomonas, Terrimonas and Sphingomonas.

Fungi phyla Ascomycota (genera: Pilidium, Dendroclathra, Rosellinia and Hypomyces) and Basidiomycota (genera: Coprinellus and Coprinus) were confirmed.

This study is the first to identify specific As- and TI- tolerant fungal (Pilidium, Cladophialophora, Neobulgaria, and Mycena) and bacterial genera (Trichococcus, Devosia, Litorilinea, and Gimesia).

ICP-OES analysis of the samples detected 25 elements and high concentration of As and TI.

Keywords: Allchar, mine, bacteria, fungi, diversity, lorandite, thallium.



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