



BOOK OF ABSTRACTS

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15-16TH JUNE 2022. SKOPJE, REPUBLIC OF NORTH MACEDONIA



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Contents

Modelling of potential distribution of the atlas cedar (<i>Cedrus atlantica</i> Manetti) i Morocco	
Mycodiversity in the coniferous communities of Shar Mountain - Literature revieus of macrofungi in coniferous forest on mountain Shar Planina in North Macedon	
Morphological variability of bulbs, fruits, and seeds of the genus <i>Galanthus</i> L. (Amaryllidaceae) in Serbia	. 14
First report of <i>Cryphonectria carpinicola</i> in the Balkans and <i>Cryphonectria</i> radicalis in Bulgaria	. 15
The role of Tharandt Botanic Garden in search for alternative tree species in climate change	. 16
Correct planning and proper landscape design - key to quality urban life Case Study: Macedonia Park, Skopje, North Macedonia	. 17
Genetic differentiation of hungarian oak (<i>Quercus frainetto</i> Ten.) Populations ir Serbia based on morphological characteristics of acorns and seedlings	
Prevention of Waste-Contaminated Forests by ICT, standards and legislation	. 19
Accuracy assesment of supervised classification employing the MLC and RF algorithms on Sentinel2 image for the preparation of a vegetation map in the eastern part of the Republic of North Macedonia	. 20
Data on Distribution and Habitat Preferences of Edible Truffles in the Republic North Macedonia	
Urban forestry and disaster relief: The importance of urban forestry and greene in the 1963 post-earthquake reconstruction of Skopje	
The role of urban trees in reducing land surface temperature	23
Conservation strategy targeting priority species of the Balkans: the example of the INTERREG programme CONSE-PP	
Collection of experiences from 25 years work on seed propagation of allochthonous dendroflora at the Hans Em Faculty of Forest Sciences, Landscape Architecture and Environmental Engineering in Skopje and the surrpoundings	. 26
Analysis of the influence of CoVid-19 pandemic on residents' perception on the importance of the urban greenery on quality of life	
Mapping and evaluation of cultural ecosystem services in City Park Skopje	. 29







Protected areas: History of their evolution and management practices	30
Policy support measures in forestry and nature conservation for climate change governance	
Importance of NWFPs for good forest governance	33
Intrapreneurship as a way for employee innovation behaviour in state-owned forest enterprises	34
Identifying runoff sensitivity in forests and seminatural areas; Case study: Skopska Crna Gora	35
Development of soil erosion and desertification maps of Republic of North Macedonia	36
Natural revegetation and overgrowing on forest roads in beech stands and undermountainous conditions	
Contribution of forest certification to sustainable development goals	39
Real growth rate and age structure of Brown hare (<i>Lepus Europaeus</i> P.) at the Serbian hunting grounds	40
The tradition of sustainable management in lowland forests of Croatia	41
Spatial optimization for forest planning	43
Recent changes in the population structure of <i>Cryphonectria parasitica</i> in Turke	
Detected changes of population structure within a <i>Cryphonectria parasitica</i> population at the site Osoj, North Macedonia, during a time-span of over two decades	45
The effect of temperature on the growth of the phytopathogenic fungus Ophiostoma novo-ulmi	46
Pedunculate oak (<i>Quercus robur</i> L.) phenological varieties exhibited a different response and recovery potential to extreme drought in Serbia	47
Forest resource dynamic in Republic of North Macedonia in the period from 196 to 2016	
Urban hedges–an important green element in the system of landscape design.	50
Criteria for Evaluation of Urban Green Spaces: A Case of Macedonian Cities	52
Soil erosion and Torrential Flood Prevention: Curriculum Development at the Universities of Western Balkan Countries	54
Dendroecology analysis of Macedonian pine (<i>Pinus peuce Gris</i> .) radial growth	56







Influence of elevation on some structural elements of Macedonian pine (<i>Pinus peuce</i> Gris.) stands in National Park Pelister in North Macedonia	58
Human dimension of large carnivores' management and conservation in Serbia	
Spatial structural characteristic of Macedonian pine (<i>Pinus peuce Gris.</i>) forest in National Park Pelister in North Macedonia	
Analysis of Croatian <i>Ophiostoma novo-ulmi</i> populations – insight into changes occurring over time	63
The valorization and valuation of historical park - Topcider in Belgrade	64
Current status of forest soil mapping in Serbia	65
Nationwide fuel models structure analysis using discrete ALS data in Portugal .	66
Corythucha arcuata (Say) predilection research results in 70 pedunculate oak clones (Quercus robur L.)	67
Growth of scots pine (<i>Pinus Sylvestris</i> L.) in international provenance tests in Žepče, Bosnia and Herzegovina	68
The role of site ecology into planning and designing of the conifers seed orchar in Serbia	
Diversity of pedunculate oak (<i>Quercus robur</i> L.) in Serbia based on phenotypic traits	
Possibilities of hybrid drives of forest vehicles	71
Cooperation and networking of non-wood forest products-based enterprises in the Topličko and Moravsko forest region	72
Forests in women's hands: Activities on the development of the National action plan for Serbia	
In situ conservation of rare and endangered forest tree species at the locality "Jelovarnik" - National Park "Kopaonik" in Serbia	75
Influence of tree crown status on the growth of common beech (<i>Fagus sylvatica</i> L.) in Croatia	
Interrelations of various tree vitality indicators on a common beech (<i>Fagus sylvatica</i> L.) plot	77
Conservation of forest fruit trees gene pool in the protected area "Košutnjak forest" (Serbia)	78
Climate changes' impact on management and activities in protected areas: Managers' and Stakeholders' attitudes	79





The impact of pest insects and fungal diseases on the dieback of pine fo	rests in
the R. of N. Macedonia	81
Bioecological characteristics of two mass outbreaks of Altica Querceorun	<i>n</i> foudr.
in the R of N. Macedonia	82
Effect of Wildfire on Forest Soils in Maleshevo-Pijanec Region	83



Analysis of Croatian *Ophiostoma novo-ulmi* populations – insight into changes occurring over time

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Abstract:

Ophiostoma novo-ulmi is a phytopathogenic ascomycete fungus that causes Dutch elm disease (DED) – a highly destructive disease responsible for killing numerous North American and European elms (*Ulmus* ssp.). DED spreading began in the early 20th century and it was originally caused by Ophiostoma ulmi, while O. novo-ulmi emerged in the mid-20th century, initiated an even more destructive DED pandemic, and eventually became the main DED pathogen. Two O. novo-ulmi subspecies (O. novo-ulmi ssp. novo-ulmi and ssp. americana) are widespread throughout Europe and their hybrids are frequently found in areas of overlapping distribution. Following the DED survey across Croatia conducted in 2012, elms were resampled in selected Croatian forests (sites Nova Kapela and Jastrebarsko) in 2021. An increase in disease incidence at site Nova Kapela was observed in 2021 compared to 2012, while at site Jastrebarsko a decrease in disease incidence was detected. Molecular characterization of O. novo-ulmi isolates based on analysis of the col1 and cu gene sequences indicated that in 2021, similar to 2012, the O. novo-ulmi ssp. novo-ulmi is more frequent compared to O. novo-ulmi ssp. americana which was only detected at site Nova Kapela in 2021. O. novo-ulmi ssp. novo-ulmi x ssp. americana hybrids were found at both sites in both years of investigation. Obtained results point out that variations in disease incidence do occur over time, but the dominancy of O. novoulmi ssp. novo-ulmi and hybrids over O. novo-ulmi ssp. americana is retained in investigated sites.

Keywords: Dutch elm disease; elms; temporal variability



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