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## Microscopic fungi isolated from the Domica Cave system (Slovak Karst National Park, Slovakia). A review

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### Abstract:

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A broad spectrum, total of 195 microfungus taxa, were isolated from various cave substrates (cave air, cave sediments, bat droppings and/or guano, earthworm casts, isopods and diplopods faeces, mammalian dung, cadavers, vermiculations, insect bodies, plant material, etc.) from the cave system of the Domica Cave (Slovak Karst National Park, Slovakia) using dilution, direct and gravity settling culture plate methods and several isolation media. *Penicillium glandicola*, *Trichoderma polysporum*, *Oidiodendron cerealis*, *Mucor* spp., *Talaromyces flavus* and species of the genus *Doratomyces* were isolated frequently during our study. Estimated microfungus species diversity was compared with literature records from the same substrates published in the past.

**Keywords:** Domica Cave system, microfungi, air, sediments, bat guano, invertebrate traces, dung, vermiculations, cadavers

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### INTRODUCTION

Microscopic fungi are an important part of cave microflora and occur in various substrates in caves, such as cave sediments, vermiculations, bat droppings and/or guano, decaying organic material, etc. Their widespread distribution contributes to their important role in the feeding strategies of cave fauna.

A study of microscopic fungi in the Domica Cave System (Slovak Karst National Park, Slovakia) was started in 2002 within the frame of the project studying cave microflora and fauna and their feeding preferences and it was subsequently carried out each spring and autumn. The first isolation was performed only on the Domica Cave (a show cave). Later, parts not open to the public were sampled as well, i.e. the chasm-like Čertova diera Cave, the Dlhá Chodba Cave and its part in Hungary with the aim to obtain a broad spectrum of microscopic fungi which can represent potential food for cave mycophagous invertebrates. Partial results of these samplings were presented at international conferences and workshops or published in scientific journals (Nováková, 2004a, b, 2005, 2006, 2008, Elhottová et al., 2003, 2004, Nováková et al., 2005, 2008, Lukešová & Nováková, 2008, Šustr et al., 2005).

The goal of this paper is to form a view of microscopic fungi isolated from different substrates and parts of the Domica Cave system and a comparison of

the obtained microfungus spectrum with records of previously published data from the Baradla Cave and other caves in the world.

### DESCRIPTION OF STUDIED CAVES

The Domica Cave system is located on the southwestern edge of the Silická Plateau in the Slovak Karst National Park, close to the state border with Hungary. Geographic coordinates of the Domica Cave entrance are 48°28'43" N and 20°28'22" E, and its entrance is counted in the land-register of the village Kečovo 10 km south of the town Plešivec, at the southern foothill of Domica Hill, 339 m a.s.l. The Domica Cave is connected with the chasm-like Čertova diera Cave (the Devil's Pit Cave) and together with the Stará Domica Cave (the Old Domica Cave) and the Dlhá Chodba Cave (Long Passage Cave - a corridor between Domica and Baradla caves) they reach a length of 5,358 m. They also form one generic unit with the Baradla Cave in Hungary with a total length of about 25 km, from which almost one quarter is in Slovak territory. Currently, the show cave is 1,315 m. The cave was formed in the Middle Triassic by pale Wetterstein limestones of the Silica Nappe along the tectonic faults by corrosive and erosive activities of the Styx stream and the Domický Brook and smaller underground tributaries draining water mainly from the non-karst part of the basin. Air temperature ranges from 10.2 to 11.4 °C and relative humidity from 95 to 98 %. The cave system is an important **wintering place** for bats. Very numerous colonies of the Schreiber's Bats (*Miniopterus schreibersii*)

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hibernated in the Čertova diera Cave in the past, and also dominant is the Mediterranean Horseshoe Bat (*Rhinolophus euryale*), which forms a unique colony of 1,200 – 1,400 members in the Domica Cave. Thick layers of bat excrements – or guano - can be found in some places in this cave system (Anonymous, 2006; Bella, 1997; Droppa, 1961; Kladiva, 1995-2008).

### MATERIAL AND METHODS

Samples for microfungal isolation were collected from various sites from the Domica Cave (both show cave tour and diversionary were sampled, DC), the Čertova diera Cave (CD) and the Dlhá Chodba Cave (DCH), i.e. from lighted and not lighted parts of the Domica Cave system (Figure 1).

Samples of cave (cave sediment, bat droppings and guano, earthworm casts, isopods and diplopods faeces, mammalian dung, frog and bat cadavers, bones, vermiculations, insects and other organic material) were collected aseptically in the spring and in the autumn of 2002 - 2007 into sterile plastic bags, vessels or microtubes and they were kept cold during the transport to the laboratory.

Microscopic fungi were isolated in situ (mainly visible colonies of the order Mucorales on bat droppings and/or guano and other traces) or immediately after return to the laboratory (i.e., most five days after sampling) either directly by transferring a small amount of collected material into Petri dishes, or using the dilution plate method (Garrett, 1981). Martin's soil extract agar, Sabouraud's agar and beer wort agar (all with rose bengal and chloramphenicol for the suppression of bacterial growth – these isolation media were found optimal for microfungal isolation from cave environment, while pure isolation

media prepared e.g. from cave sediment extract were unsuitable considering low possibility of isolation) were used as isolation media (Kreisel & Schauer, 1987). Air microfungi were isolated using the gravity settling culture plate method (Buttner & Stetzenbach, 1991) and Sabouraud's agar as the isolation medium. Petri dishes were incubated in the dark at 25 °C for 7-10 days. Additionally, soil samples from above-ground environment were analysed and microfungal isolations from the outdoor air were carried out, too.

Identifications were carried out according to micro- and macromorphological characteristics using special identification media (MEA, CYA, Czapek-Dox agar, PDA, CA, OA, etc.) for all isolated micromycetes. Special taxonomic literature and compendia (Domsch et al., 1980, 2007; de Hoog et al., 2000) were used for the determination of microfungi. Taxonomic names are used according to Domsch et al. (2007).

### RESULTS AND DISCUSSION

A total of 195 microfungal taxa (Zygomycetes, Ascomycetes and mitosporic fungi) belonging to 73 microfungal genera have been isolated during the study in the Domica Cave system (Table 1), and mainly saprotrophic filamentous microfungi were studied. The highest number of microfungal taxa (92) was found in samples of bat droppings and guano. Bat droppings and guano were collected in three parts of the Domica Cave system (the Čertova diera, Domica and Dlhá Chodba caves) – the guano heaps in the Domica Cave (Figure 2) and in the Čertova diera Cave make up the largest guano heaps in Central European caves. 15 microfungal taxa were isolated from Domica and Čertova diera caves, 11 taxa were obtained only from the Čertova diera Cave, 52 taxa were isolated only

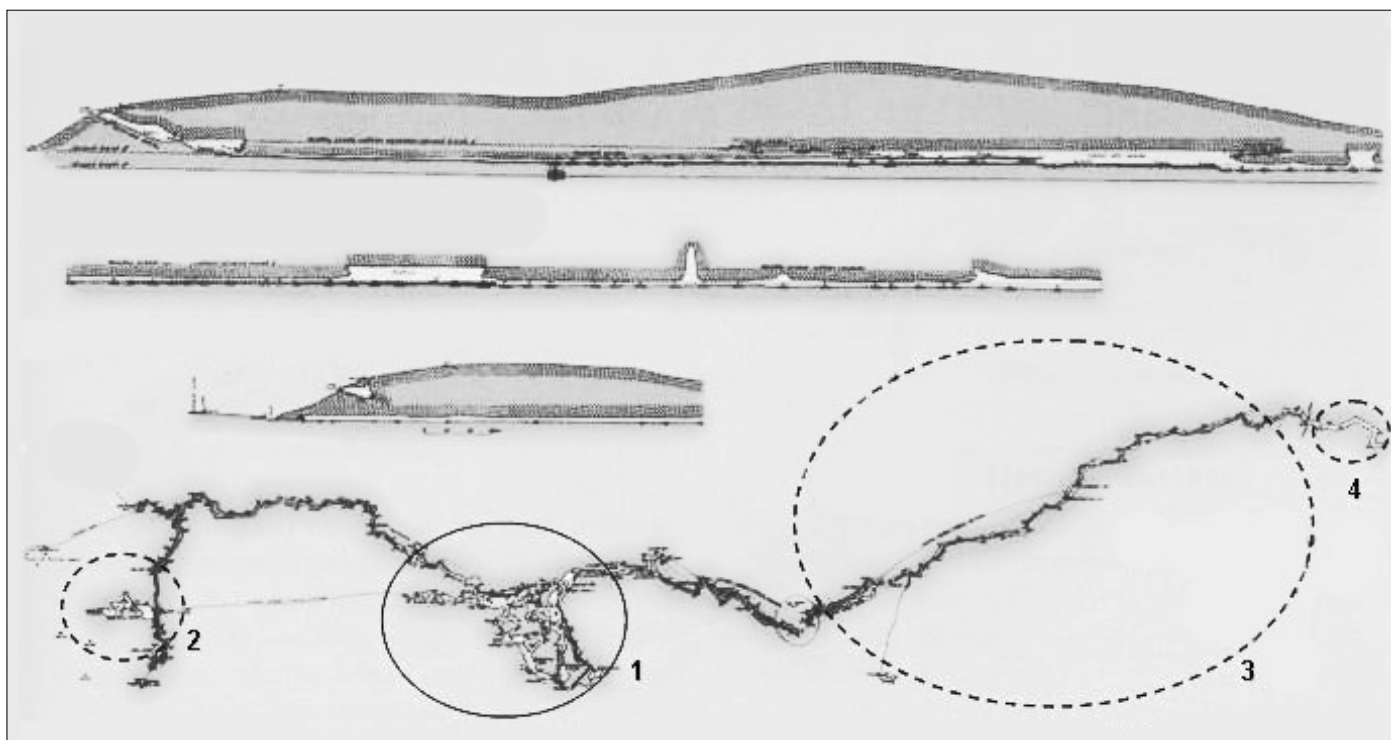


Fig. 1. A map of the Domica Cave system created by A. Droppa and A. Chovan – adapted according to Jakál (2005). Continuous line – show cave, broken line - non-tourist areas (1 – the Domica Cave, 2 – the Čertova diera Cave, 3 – the Dlhá Chodba Cave, 4 - Hungarian part of the Dlhá Chodba Cave).

Tab. 1. A view of microfungus taxa isolated from the cave sediment air, bat droppings and/or guano, mammalian dung, earthworm casts, isopod and diplopods faeces, vermiculations, cadavers, and other substrates. 1 – the Domica Cave, 2 – the Čertova diera Cave, 3 – the Dlhá Chodba Cave, 4 - Hungarian part of the Dlhá Chodba Cave; frequent isolations are in bold.

microfungal taxa	sediment	air	bat guano	mammalian dung	earthworm casts	isopod diplopod faeces	vermiculations	cadavers	other substrates
<i>Zygomycetes:</i>									
<i>Absidia cylindrospora</i> var. <i>cylindrospora</i>			2		1				
<i>Absidia cylindrospora</i> var. <i>nigra</i>	1								
<i>Absidia glauca</i>			1						3
<i>Absidia spinosa</i>	1								
<i>Circinella</i> sp.		3							
<i>Coemansia aciculifera</i>					4				
<i>Mortierella</i> spp.	1		1,3		1	1		1,3	1
<i>Mucor circinelloides</i> f. <i>circinelloides</i>		3	1						
<i>Mucor dimorphosporus</i> f. <i>dimorphosporus</i>	3	2	1						3
<i>Mucor dimorphosporus</i> f. <i>sphaerosporus</i>		2	1, 2						3
<i>Mucor hiemalis</i> f. <i>corticola</i>			2						
<i>Mucor hiemalis</i> f. <i>hiemalis</i>		1,2	1,2						
<i>Mucor hiemalis</i> f. <i>silvaticus</i>		1,2	1		1			1	
? <i>Mucor mucedo</i>			1,2,3						
<i>Mucor ramosissimus</i>		1							
<i>Mucor</i> spp.			1	1					1
<i>Rhizopus arrhizus</i>		1,2,3							
<i>Rhizopus stolonifer</i> var. <i>stolonifer</i>	2	2	1						
<i>mitosporic fungi:</i>									
<i>Acremonium bactrocephalum</i>						1			
<i>Acremonium berkeleyanum</i>		1	1,2			1			
<i>Acremonium charticola</i>			1						
<i>Acremonium murorum</i>	1,2		1		1,2				1
<i>Acremonium polychromum</i>	1								
<i>Acremonium strictum</i>					1				
<i>Acremonium</i> sp.	1,3	2			1,3				
<i>Acrodontium</i> sp.		1							
<i>Acrostalagmus luteoalbus</i>	1,2,3								
<i>Alternaria alternata</i>		1,2,3	1						1
<i>Arthrinium arundinis</i>		1	1,2		1				
<i>Arthrinium phaeospermum</i>	1,3				2				
<i>Aspergillus candidus</i>		1				1			
<i>Aspergillus clavatus</i>			1						
<i>Aspergillus flavus</i> group	3								
<i>Aspergillus fumigatus</i>	3	1,2	1,2		1,2				
<i>Aspergillus niger</i> group			2						
<i>Aspergillus phoenicis</i>	1				1				1
<i>Aspergillus puniceus</i>					2				
<i>Aspergillus ustus</i>		1							1
<i>Aspergillus versicolor</i>		1,2				1			
<i>Aspergillus</i> sp.	1, 2	1,3	2			1	1		1
<i>Beauveria bassiana</i>	1,2,3	1,2	1,2						1
<i>Beauveria brongniartii</i>		1							1
<i>Botryosporium longibrachiatum</i>								1,4	
<i>Botryotrichum piluliferum</i>	1		2						
<i>Botrytis cinerea</i>		2			2,3				

<i>Chloridium virescens</i> var. <i>candigerum</i>					<b>1</b>				
<i>Chrysosporium speluncarum</i>			2						
<i>Chrysosporium</i> sp.	1, 4	1	<b>1,2</b>		<b>1,2</b>		3		
<i>Cladosporium cladosporioides</i>	1	<b>1,2,3</b>							
<i>Cladosporium herbarum</i>	1	<b>1,2,3</b>							
<i>Cladosporium</i> cf. <i>elatum</i>		1,2							
<i>Cladosporium sphaerospermum</i>		<b>1,2</b>							
<i>Cladosporium tenuissimum</i>		<b>1</b>							
<i>Cladophialophora</i> sp.			1		<b>1</b>				
<i>Clonostachys candelabra</i>		2							
<i>Clonostachys rosea</i> f. <i>rosea</i>	1		<b>1,2</b>		<b>1</b>	<b>3</b>			
<i>Clonostachys</i> sp.	3								
<i>Cylindrocarpon destructans</i>					1				
<i>Cylindrocarpon</i> sp.					1	3			
<i>Doratomyces microsporus</i>	<b>1</b>					3			
<i>Doratomyces nanus</i>			<b>1,2</b>		2	3			
<i>Doratomyces stemonitis</i>	<b>1,2</b>	4	<b>1,2</b>						
<i>Echinobotryum</i> state of <i>Doratomyces stemonitis</i>	<b>3</b>		<b>1,2</b>			3			
<i>Emericellopsis terricola</i>	1								
<i>Engyodontium album</i>	2	1,2							
<i>Fusarium solani</i>			1						
<i>Fusarium sporotrichoides</i>	1								
<i>Fusarium</i> spp.	1,4	2,3	<b>1</b>		<b>1</b>		1		
<i>Geomyces pannorum</i> var. <i>pannorum</i>	<b>1</b>	<b>1</b>	<b>1,2</b>		1,2	3			
<i>Geotrichum candidum</i>			<b>1</b>				<b>3</b>		1
<i>Hormiactis</i> sp.			1		1				
<i>Humicola fuscoatra</i>	1, 3		1						1
<i>Humicola grisea</i>	1,2		2		1,2				
<i>Humicola</i> sp.					1				
<i>Hypocrea</i> sp.			1						
<i>Isaria farinosa</i>	1,2,3	<b>2,3</b>	1,2		1				1
<i>Isaria fumosorosea</i>	1		2		2				
<i>Lecanicillium muscarium</i>	<b>1</b>		<b>1</b>						1
<i>Lecanicillium psalliotae</i>		2							
<i>Malbranchea</i> sp.	1,4	2	1		4				
<i>Mammaria echinobotryoides</i>	1				2				
<i>Metarhizium anisopliae</i>								<b>3</b>	
? <i>Myceliophthora velerea</i>		1							
<i>Myrothecium roridum</i>	3		2		3,4				
<i>Myrothecium verrucaria</i>	2,1								
<i>Myxotrichum deflexum</i>	<b>1,3</b>		2		<b>1</b>		1		
<i>Ochroconis</i> ( <i>Scolecobasidium</i> ) <i>tschawytschae</i>		3							
<i>Oidiodendron cerealis</i>	<b>1,3</b>	<b>3</b>	<b>1</b>		<b>1,3</b>		1,4		1,3
<i>Oidiodendron citrinum</i>			1						
<i>Oidiodendron griseum</i>	<b>1,2</b>		<b>1</b>				3		
<i>Oidiodendron tenuissimum</i>	1								
<i>Oidiodendron</i> sp.	<b>3</b>		<b>1</b>						
<i>Paecilium lilacinum</i>	<b>1,3</b>	1	1	1	1,3	1			
<i>Paecilomyces marquandii</i>	<b>1</b>		3		1	3			
<i>Paecilomyces variotii</i>	1,2,3		1,2		<b>1</b>				
<i>Paecilomyces</i> sp. ( <i>white</i> )	3								
<i>Paranomurea carnea</i>	<b>1,3</b>	1,2	<b>1,3</b>		<b>1,2</b>		1,3		1

<i>Penicillium atramentosum</i>					2				
<i>Penicillium aurantiogriseum</i>		1	1		1				
<i>Penicillium chrysogenum</i>		2	1		1				
<i>Penicillium citrinum</i>	1	1	1,2		2				
<i>Penicillium commune</i>		1	1						
<i>Penicillium corylophilum</i>	1	1	2						
<i>Penicillium daleae</i>			1				1		
<i>Penicillium decumbens</i>		1							
<i>Penicillium echinulatum</i>			1,2						
<i>Penicillium expansum</i>			1,2		1				3
<i>Penicillium glabrum</i>	1		1						
<i>Penicillium glandicola</i>			1,2,3	1	1,2				1
<i>Penicillium hirsutum</i>			1						
<i>Penicillium hordei</i>			1						
<i>Penicillium janczewskii</i>		1			1,2				
<i>Penicillium janthinellum</i>			1						
<i>Penicillium melanoconidium</i>			1						
<i>Penicillium melinii</i>		1			1				
<i>Penicillium minioluteum</i>			1,2		1	1			
<i>Penicillium paxilli</i>					1				
<i>Penicillium pinophilum</i>		1	1		1				
<i>Penicillium purpurescens</i>					1				
<i>Penicillium purpurogenum</i>	1								
<i>Penicillium restrictum</i>		1							
<i>Penicillium roqueforti</i>			1						
<i>Penicillium rugulosum</i>	1				1,3	3			
<i>Penicillium sacculum</i>	1,2				1				
<i>Penicillium scabrosum</i>	1								
<i>Penicillium solitum</i>									1
<i>Penicillium thomii</i>					1				
<i>Penicillium variabile</i>		1	1		1		1		
<i>Penicillium viridicatum</i>		1							
<i>Penicillium vulpinum</i>		2	1						
<i>Penicillium waksmanii</i>					3				
<i>Penicillium sp.</i>		1	1	1	1		1		1
<i>Periconia cf. macrospinoso</i>					1				
<i>Phialophora sp.</i>		1							
<i>Phoma eupyrena</i>	1,3				3				
<i>Phoma lingam</i>		2							1
<i>Phoma sp.</i>		1			1				1
<i>Pithomyces chartarum</i>		1,2							
<i>Pochonia chlamydosporia var. catenulata</i>	3,4,1	1	1,2		2				
<i>Pochonia chlamydosporia var. chlamydosporia</i>	3								
<i>Ramichloridium sp.</i>	1								
<i>Scopulariopsis brumptii</i>					1,3	1			
<i>Scopulariopsis chartarum</i>	1								
<i>Scopulariopsis croci</i>			1						
<i>Scytalidium lignicola</i>	1,3	1			1				
<i>Simplicillium lamellicola</i>	1		1,2						
<i>Stachybotrys chartarum</i>		2							
<i>Stachybotrys cylindrospora</i>	1								
<i>Tetracosporium paxianum</i>	1,2					2			

<i>Tolypocladium niveum</i>			1		2					
<i>Tolypocladium cylindrosporum</i>	1									
<i>Trichocladium opacum</i>	1									
<i>Trichoderma atroviride</i>	1	1	1			1				
<i>Trichoderma hamatum</i>	1		1			1			1	
<i>Trichoderma harzianum</i>									1	
<i>Trichoderma koningii</i>									1	
<i>Trichoderma polysporum</i>	1,3	1,2	1		1,3,4	1,2	1,4	1	1	
<i>Trichoderma state of Hypocrea stellata</i>					3					
<i>Trichoderma sp.</i>	1	1	1		1	1		1		
? <i>Trichophyton erinacei</i>			1							
<i>Trichophyton sp.</i>		1								
<i>Truncatella angustata</i>	1,4									
<i>Ulocladium atrum</i>		1								
<i>Ulocladium chartarum</i>		2,3								
<i>Ulocladium oudemansii</i>		2								
<i>Ulocladium sp.</i>		3								
<i>Verticillium sp.</i>	3	1	1,2,3							
<i>Walemia sebi</i>					3					
<i>Wardomyces anomalus</i>					3					
<i>Wardomyces sp.</i>						3				
<b>Ascomycetes:</b>										
<i>Chaetomium crispatum</i>					2					
<i>Chaetomium funicola</i>	1					1				
<i>Chaetomium globosum</i>					1					
<i>Chaetomium indicum</i>	1,3		1		1		1			
<i>Chaetomium spinosum</i>			1		1					
<i>Chaetomium sp.</i>	1,3				1,3,4	1	1			
<i>Emericella nidulans</i>			1							
? <i>Eupenicillium javanicum</i>					1					
<i>Eupenicillium sp.</i>					1					
<i>Eurotium amstelodami</i>			1							
<i>Eurotium sp.</i>	1,4									
<i>Gymnoascus reessii</i>	1,3		1				1			
<i>Pidoplitchoviella terricola</i>					1					
<i>Talaromyces flavus</i>	1,3,4		1,2		1,3,4	3	1		1	
<i>Talaromyces wortmannii</i>					1,2,3			3	3	
<i>Talaromyces sp.</i>	1,3				1,2,3,4		1	4	1	
<i>Talaromyces (white)</i>	1,4				1					
<i>Thielavia hyrcaniae</i>					1					
<i>unidentified fungi:</i>										
sterile white mycelium	1,4	1,2	1		1	1				
sterile yellow mycelium	3		1		1,3,4	3		3	1	
sterile beige mycelium							1			
sterile pink mycelium	1	2								
dark sterile mycelium	1		1		1					
undetermined species os Ascomycetes	3		1		1,4				3	
undetermined species of Dematiaceae	3				1,3	4	1			
undetermined species of Sphaeropsidales					3	3				
Total numbers of isolated taxa	195	86	73	92	4	86	30	19	9	33



Fig. 2. Guano heap in the Palmy Grove Dome in the Domica Cave - the greatest guano heap in Central Europe caves (depth of 103 cm, heap base of 440 x 350 cm).

from the Domica Cave and six microfungi taxa were found in bat droppings and/or guano samples from all parts of the Domica Cave system. Several microfungi taxa were isolated from this material repeatedly, e.g. *Absidia glauca*, *Acremonium berkeleyanum*, *Alternaria alternata*, *Aspergillus clavatus*, *A. fumigatus*, *Chrysosporium* sp., *Clonostachys rosea* f. *rosea*, *Geomyces pannorum* var. *pannorum*, *Lecanicillium muscarium*, *Mucor* spp., *Oidiendendron cerealis*, *O. griseum*, *Oidiendendron* sp., *Doratomyces stemonitis* and its co-anamorph *Echinobotryum* (Figure 3), *Trichoderma polysporum* (Figure 4), *Paranomuraea carnea*, *Penicillium citrinum*, *P. glandicola* (Figure 5), *Rhizopus arrhizus*, and *Ulocladium* sp. *Oidiendendron cerealis*, *Doratomyces stemonitis* with co-anamorph *Echinobotryum*, and *Trichoderma polysporum* were the most frequently isolated species, and may represent typical microfungi species on bat traces. Species of the genus *Mucor* which create visible colonies on bat droppings and on the surface of guano heaps were also isolated with high frequency using direct isolation in situ. In all, about 30 species of this genus differing in micro- and macromorphological properties were recognised from this substrate – unfortunately they were not determined because a monograph with a modern concept of this genus was not available at that time. Eleven microfungi species were found only from bat droppings and/or guano and were not found in other studied materials (see Table 1). White and yellowish pustules were found on bat droppings in the Domica Cave system as well as in other caves in the NP Slovak Karst, but the isolation was unsuccessful. Finally, *Trichoderma polysporum*, one of the white-coloured *Trichoderma* species, was determined from white colonies. In microscopic slides prepared from yellowish microfungi colonies on bat droppings, tuberculate conidia resembling anamorph of *Ajellomyces capsulatus* were observed. Considering the successful isolation of this fungus from bat guano in 2006 and consequential analyses including molecular techniques, novel *Chrysosporium* species, *C. speluncarum*, was described (Nováková & Kolařík, in press).

A broad spectrum of microfungi species are known from previous studies of microfungi from bat droppings and guano; e.g., Larcher et al. (2003) revealed the great fungal diversity from 82 studied samples of bat guano from caves of Western France, including pathogenic yeasts, eight species of keratinophilic fungi, and one onygenalean fungus highly resembling *Histoplasma capsulatum*. *Trichosporon laibachii*, *T. porosum* and seven new *Trichosporon* species, eight ascomycetous yeasts and one basidiomycetous yeast were reported from bat guano samples from bat-inhabited caves in Japan (Sugita et al., 2005). Ulloa et al. (2006) isolated four ascomycetal species from bat guano samples collected in several Mexican caves (*Aphanoascus fulvescens*, *Gymnascella citrina*, *Gymnoascus dankaliensis*, *Chaetomidium fimeti*), six mitosporic fungi (*Aspergillus flavofurcatis*, *A. terreus*, *A. terreus* var. *aureus*, *Penicillium* spp., *Malbranchea aurantiaca*, and *Sporothrix* sp.), and five yeasts (*Rhodotorula* spp., *Candida catenulata*, *C. cifferii*, *C. famata* var. *flareri*, and *C. guilliermondii* var. *guilliermondii*). Several morphological types of yeasts were also found between isolated filamentous microfungi, but they were not isolated in this study.

Important differences in species diversity and numbers of isolated microfungi species from bat droppings and/or guano were found between our results and literature records. Low microfungi numbers and pure spectrum which are reported in some above mentioned papers quite agree with using isolation techniques and mainly with short-time of investigation.

Microfungi studies in the Baradla Cave, a part of the Domica-Baradla Cave System in Hungary, performed in the 1960s by Zeller (1962, 1966, 1967, 1968, 1970) targeted microfungi isolation largely using the To-Ka-Va baiting method, and some microfungi species were also isolated using the dilution plate method. *Arthoderma quadrifidum*, *A. currei*, *A. tuberculatum*, *Trichophyton terrestre*, *Myxotrichum chartarum*, *M. deflexum*, *Chrysosporium evolceanui*, *Circinella simplex*, *Rhizopus arrhizus*, *R. nigricans*, *R. delemar*, and *Absidia corymbifera* were isolated from bat guano in these studies. *Myxotrichum deflexum* was also isolated from guano samples from the Čertova diera Cave and repeatedly from cave sediment (DC, DCH) and earthworm casts (DC), and in one case this species was found from vermiculations (DC), too (see Table 1).

Some differences in microfungi diversity in cave sediments and earthworm casts (Figure 6a, b) were found after six years of investigations, but in total 86 taxa were isolated from both substrates even though a higher number of cave sediments were analysed compared to the number of sampled earthworm casts, although species richness always seemed to be higher in earthworm casts. The greater part of the isolated microfungi taxa was found in both substrates, some of which were isolated repeatedly, but several taxa were recorded only from casts (e.g., *Chaetomium crispatum*, *Doratomyces nanus*, *Tolypocladium niveum*, *Talaromyces wortmannii*,

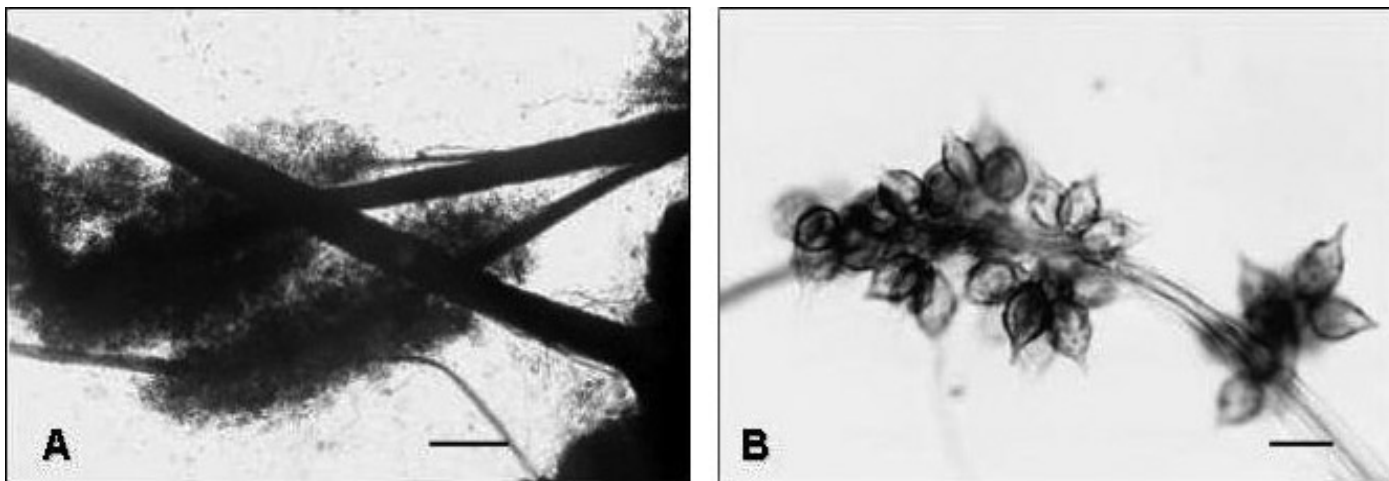


Fig. 3. *Doratomyces stemonitis* and its co-anamorph *Echinobotryum* - A – synnemata of *D. stemonitis* (bar = 20 µm); B – *Echinobotryum* type aleurioconidia (bar = 10 µm).

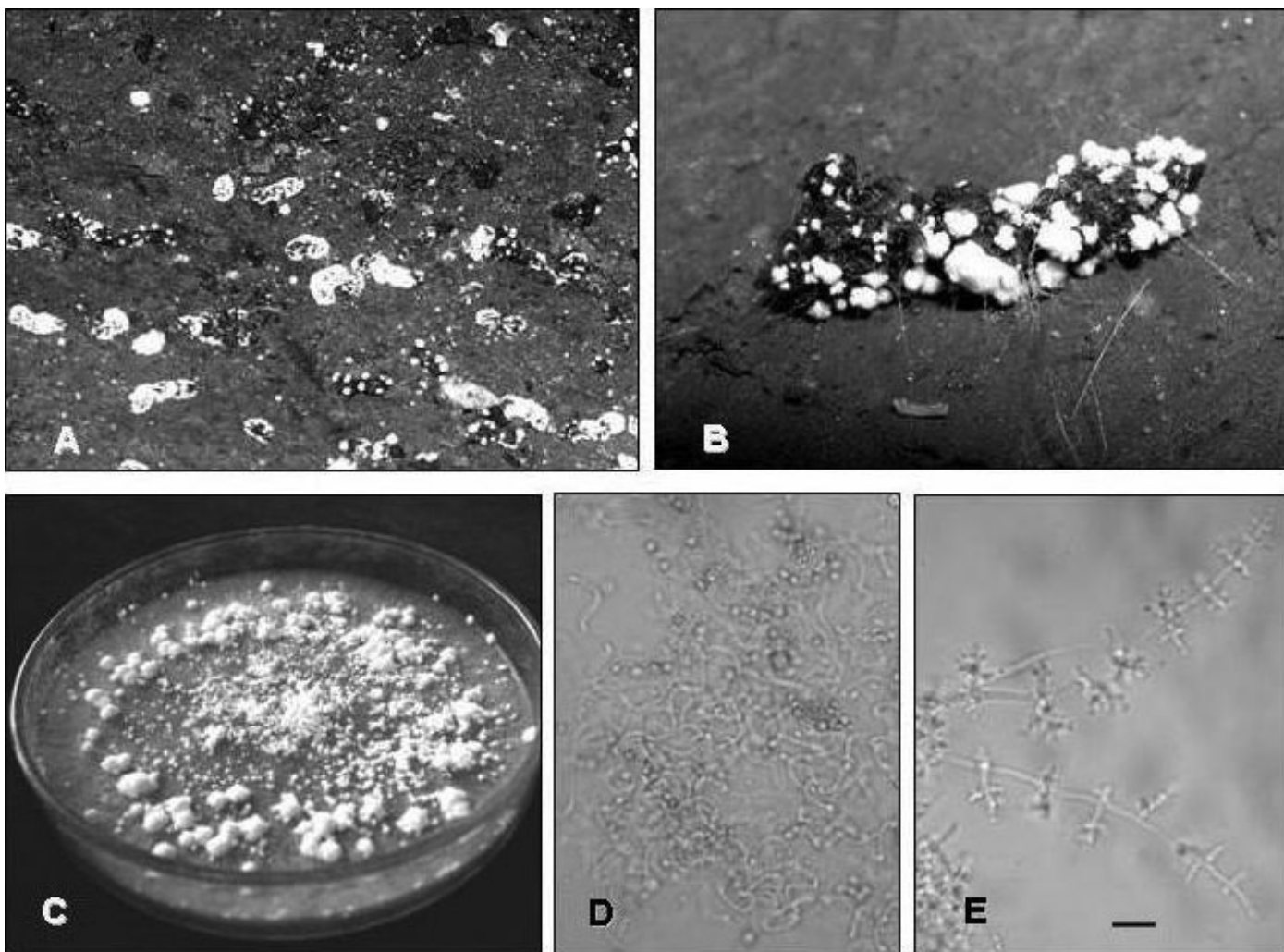


Fig. 4. A – Bat droppings with white microfungal colonies; B – bat dropping with white pustules of *Trichoderma polysporum*, photo J. Stankovič; C – *T. polysporum*, 10 days colony on malt extract agar (MEA); D, E – *T. polysporum* – typical conidiophores and sterile apical elongations (bar = 10 µm).

*Walleimia sebi*, and *Wardomyces anomalus*) or cave sediments (*Arthrinium phaeospermum*, *Aspergillus fumigatus*, *Eurotium* sp., *Doratomyces microsporus*, *D. stemonitis* and its co-anamorph *Echinobotryum*, and *Emericellopsis terricola*). *Talaromyces flavus* (Figure 6c) was found in both of these substrates, but more frequently it was isolated from earthworm casts. Rutherford & Huang (1994) discussed the isolated microfungal spectrum from remote sediments in West

Virginia caves together with results of several works dealing with microfungi occurrence in sediments (and some other substrates such as dung, active cave formations, living organisms, wood and plant debris etc.) of different caves and soils in above-ground environments from around the world. They isolated 35 fungal taxa and Mycelia sterilia were found to be the most prominent forms, followed in decreasing order by *Aspergillus aureolatus*, *Byssoschlamys fulva*,



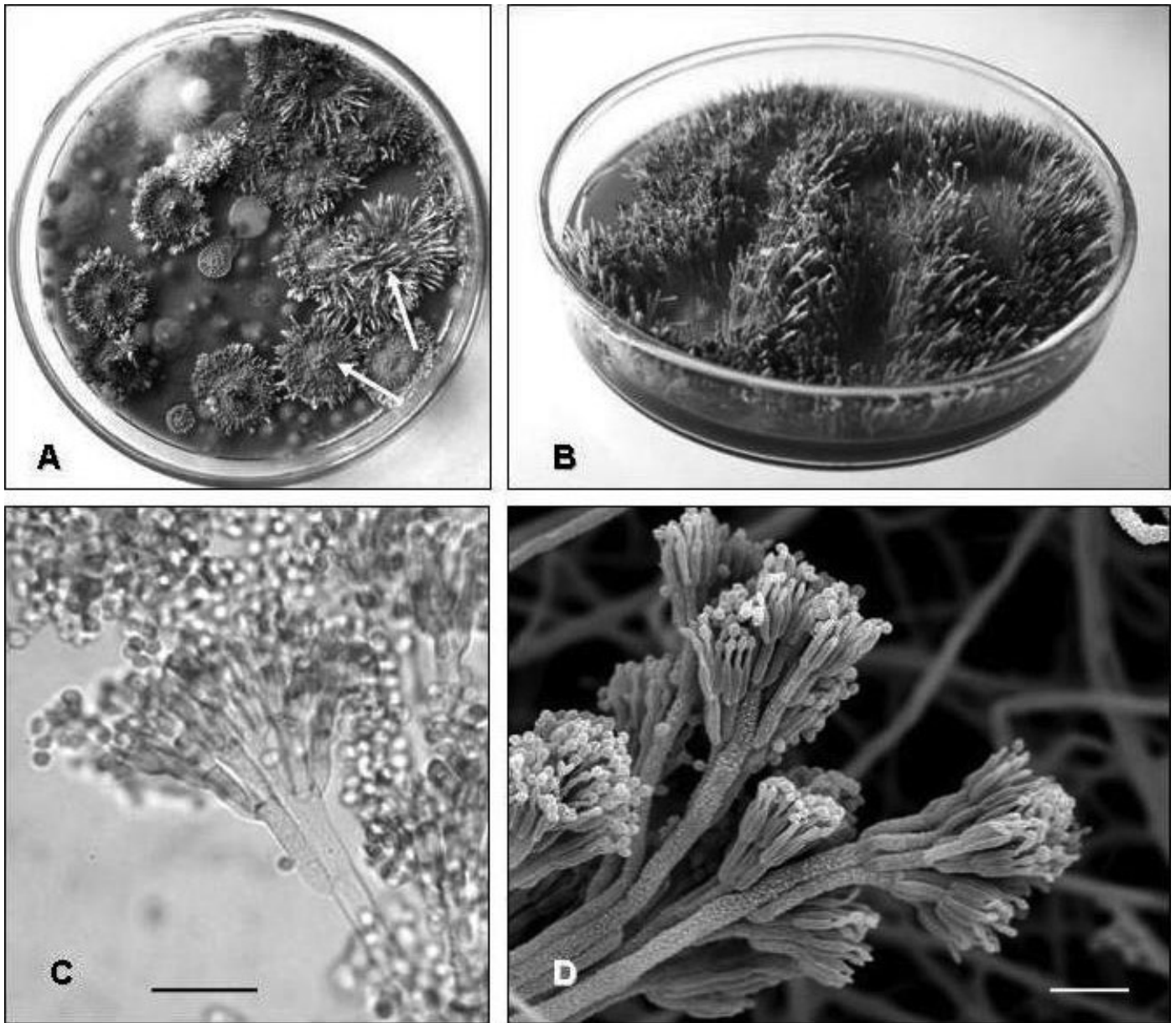


Fig. 5. A – Petri dish with microfungal colonies isolated from bat guano (Čertova diera Cave), arrows show colonies of *Penicillium glandicola*; B *P. glandicola*, 7 days synnematus colony on MEA; C – conidiophore with conidia, x 900; D – conidiophores, SEM, x 2500 (bar = 10 µm).

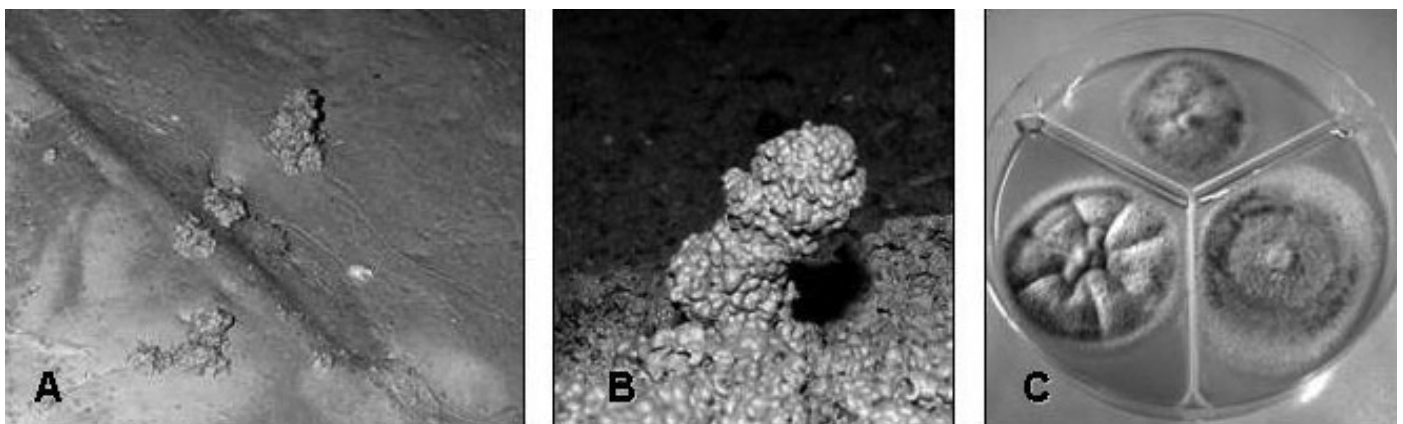


Fig. 6. A – Earthworm casts on Styx stream bank in the Domica Cave; B – funny shaped earthworm cast near Japanese Teahouse in the Domica Cave; C – *Talaromyces flavus*, 10 days colonies on Czapek-Dox agar, CYA and MEA.

*Penicillium steckii*, *Gliocladium roseum*, *Paecilomyces variotii*, *Mortierella alpina*, *Aspergillus caespitosus*, and *Fusarium oxysporum*. *Aspergillus fumigatus* was also reported from cave sediments by Griffiths (1979). A broad spectrum of micromycetes was found from a

bat cave in the Bahamas (Orpurt, 1964), Polish caves (Nespiak, 1970), the Hungarian cave Baradla (Zeller, 1962, 1966, 1968, 1970), and “la Salle de la Verna“ in the French cave of Pierre St. Martin (Seigle-Murandi et al., 1980).

Bosák et al. (2001) reported 151 microfungal taxa from four Moravian caves (Zbrašov Aragonite Caves, Javoříčko Caves, Punkva Caves, and Amatérská Cave) – 109 taxa were isolated from cave air, 56 taxa from aragonite speleothems in the Zbrašov Aragonite Caves, 35 taxa from cave sediment in the Javoříčko Caves and 13 taxa from soda straw stalactites. A higher number of microfungal taxa was found in the touristic Punkva Caves and the most frequent fungi in this cave were *Aureobasidium pullulans*, *Chrysosporium merdarium*, *Cladosporium cladosporioides*, *C. herbarum*, *C. sphaerospermum*, *Geomyces pannorum*, *Penicillium brevicompactum*, *P. vulpinum* and basidiomycetous anamorphs, whereas four microfungal species were isolated from all studied substrates (*C. macrocarpum*, *Geomyces pannorum*, *Mortierella alpina*, and *P. aurantiogriseum*), but 89 species were isolated only from one type of the studied substrates.

A comparison of the obtained microfungal spectrum from cave sediments of the Domica Cave system with microfungi isolated from soils collected above the studied cave system shows that they have very different microfungal communities. Microfungal species commonly isolated from soils were not obtained from cave sediments.

No records about micromycete occurrence in earthworm casts were published from cave environment and the same situation applies to studies of micromycete occurrence in other invertebrate traces such as isopods or diplopods which are abundant cave-dwelling organisms. In this study, a total of 30 microfungal taxa have been found from isopod and diplopod faeces while some of these were recorded only from this substrate (e.g., *Acremonium bactrocephalum*, unidentified species of Sphaeropsidales, and *Wardomyces* sp.)

Analogically, the microfungal occurrence in cave air was also occasionally studied (Figure 7). In this study, a total of 73 microfungal taxa were estimated in cave air. In comparison with outdoor mycoflora, the microfungal diversity in cave air was rather higher, but quantitative parameters (spore number in cubic meter of air) were lower. No effect on quantitative occurrence of air microfungi caused by visitors was recorded. On the contrary, Bosák et al. (2001) compared mycobiota in four Moravian caves differing in number of visitors and reported differences in

species composition including the microfungal spectrum in cave atmosphere. Very interesting is the occurrence of several species of the genus *Aspergillus* (DC, CD), because they are treated as allergenic for humans, including *A. fumigatus*, which is a potential pathogen (Domsch et al., 2007). This species was also found in cave air in the tourist Punkva Caves (Bosák et al., 2001). Several *Cladosporium* species, the entomopathogenic species *Isaria farinosa*, *Trichoderma polysporum*, and *Ulocladium chartarum* were isolated repeatedly.

Vermiculations on cave walls and speleothems were sampled only in the last two years and while 19 microfungal taxa were found in this material, only one species, *Geotrichum candidum*, was isolated repeatedly. In the past, various theories about vermiculation origin as well as participation in the processes of development of these formations on cave walls and speleothems were published (Anelli & Graniti., 1967; Bini et al., 1978; Northup et al., 2000; Northup & Lavoie, 2001; Barton, 2006; Barton & Northup, 2007). Camassa & Febroriello (2003) speculated that *Geotrichum* sp. plays an important role in the processes of vermiculation development.

Nine microfungal taxa were isolated from mammalian cadavers and bones and 33 taxa from other materials such as dead insect bodies on cave walls in the entrance parts of caves, microfungal colonies on walls and speleothems, decaying plant debris or wood material etc. Entomopathogenic species such as *Beauveria bassiana*, *B. brongniartii*, *Isaria farinosa*, and *Lecanicillium muscarium* were found on insect bodies or on walls in the entrance parts of caves. *Aspergillus ustus*, *Phoma lingam*, and *Trichoderma harzianum* were repeatedly isolated from plant debris or decaying wood material.

## CONCLUSIONS

Our long-time studies of microscopic fungi in the Domica Cave system showed a very broad microfungal spectrum, specially in cave air, sediments, invertebrate faeces and bat droppings and/or guano. Apparently, a very specific microfungal community exists in the underground environment, particularly when there are organic matter inputs. These findings are interesting mainly because micromycetes can be a potential food source

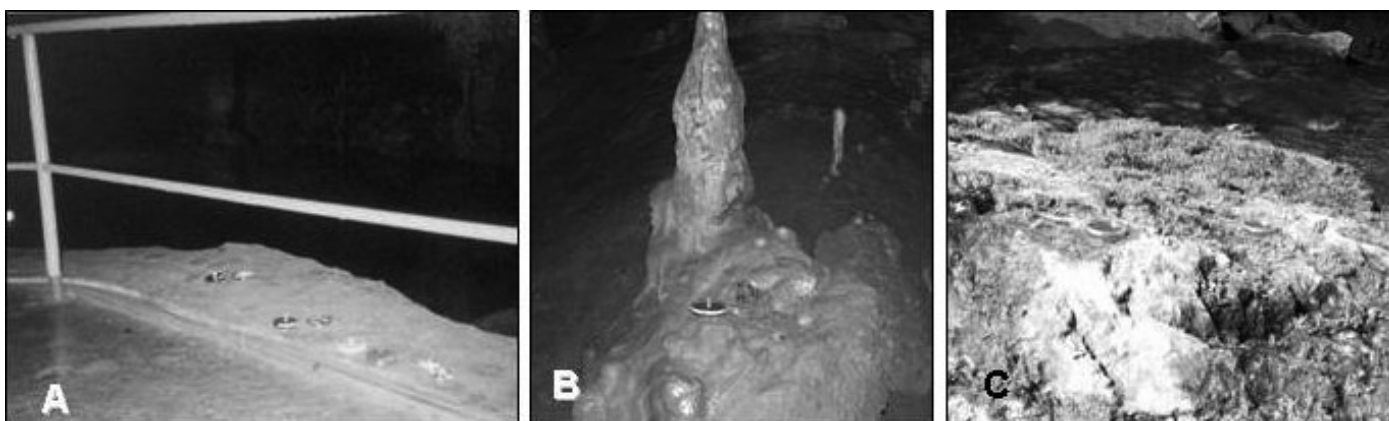


Fig. 7. Gravity settling culture plate method - isolation of air-borne micromycetes from cave air - A, B and from outdoor – C.

for cave invertebrates (together with cave bacteria and algae). *D. stemonitis* and its co-anamorph *Echinobotryum* together with *D. microsporus* and *D. nanus*, *Oidiodendron cerealis*, and *Trichoderma polysporum* were the most frequently isolated species from bat droppings/guano. *Talaromyces flavus* was frequently isolated from earthworm casts and cave sediment and *Penicillium glandicola* was found as characteristic species on marten dung and in bat guano, too. Interesting findings are also rare isolated (*Pidoplitchkoviella terricola*) or novel species (*Chrysosporium speluncarum*, sp. nov.).

It appears evident that in next studies it will be necessary to use also other isolation methods e.g., the bait technique with various type of bait, the soil washing technique and methods of molecular biology (identification of isolated strains, estimation of non-cultivated microfungi in cave substrates).

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