

Indoor Air Quality: Focus on Fungi

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Aim

To perform an indoor air quality evaluation assessing the fungal communities present in the air and surfaces of four selected Portuguese Archives. The study also includes the surface sampling of ancient documents.

Introduction

Fungi can pose a threat to both human and cultural heritage. In documents, fungi are considered responsible for the formation of foxing spots and the general degradation of written heritage. In contact with the skin or when inhaled by workers and readers, fungi may cause health problems either by toxin's emission or by the presence of fungal debris or spores. Nevertheless, the complete study of these fungal communities is still giving its first steps since only now molecular techniques are being used [1]. This project aims to assess the indoor air quality in archives, with a particular emphasis on fungal development. One of its main goals is to develop a standard procedure for the identification of the individual components of fungal communities present on the cultural written heritage stored in archives. To identify them a culture independent technique – DHPLC – is being put to use. This method allows separation of PCR products using an ion-pair reversed-phase high performance liquid chromatography and has offered exceptional advantages when it comes to the identification and isolation of microorganisms from complex microbial communities [2] [3]. The study also includes a chemical and physical parameters analysis of the environment surrounding both documents and workers.

Methods

Chemical assessment

Equipment used in the chemical evaluation of the environment. Overall, 15 rooms were analysed



Handheld 3016 IAQ
Multi Rae and Babur

Biological assessment

In the air... In the ceiling surface... In the document's surface...



Using Millipore Air Tester
20 rooms have been analysed so far



Sampling was done whenever fungal growth was visible (1 case)



A total of 15 documents have been analysed so far

Culture media: Agar, MEA and DG18



Amplification of the ITS2 and D2 genomic regions



PCR



Water activity determination
(Rotronic,TISE)

Results

Chemical results from one of the archives

Parameter	Sampling site				
	Exterior (ref. point)	A	B	C	D
O ₃ (ppm)	0.59	0.7	0.69	0.7	0.68
Formaldehyde (ppm)	0	0.003	0.002	0	0
CO (ppm)	0	1	0	0	0
VOCs (ppm)	0	0	2	2.6	0
CO ₂ (ppm)	449	604	504	471	451
Particulate matter (PM10) mg/m ³	0.042	0.129	0.047	0.033	0.147
Temp.(°C)	9.9	17	18.1	18.6	15.4
RH(%)	51.4	47.5	70.2	77.1	46.2

Parameters determined in the selected Archive. These were then analysed in terms of human health and document's conservation

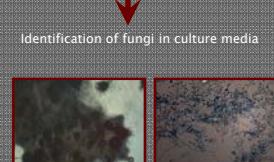
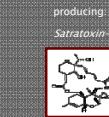
Air sampling – Analysis performed after incubation (malt extract agar, 27°C, 5-7 days)

Site	Fungi identified	CFU/m ³	Total
Exterior	<i>Aspergillus terreus</i>	4	136
	<i>Aureobasidium</i> sp.	16	
	<i>Fusarium incarnatum</i>	12	
	<i>Alternaria</i> sp.	12	
	<i>Aspergillus ochraceus</i>	4	
	<i>Ulocladium</i> sp.	4	
Office	<i>Cladosporium</i> sp.	56	96
	<i>Penicillium</i> sp.	12	
	<i>Rhodotorula</i> sp.	4	
	Yeast	12	
	<i>Aspergillus</i> sp.	8	
	<i>Penicillium</i> sp.	12	
Archive	<i>Alternaria</i> sp.	12	36
	<i>Graphium</i> sp.	12	
	<i>Cladosporium</i> sp.	24	
	<i>Paecilomyces</i> sp.	4	
	Yeast	24	
	<i>Aspergillus ochraceus</i>	8	

Colony forming units (CFU) determined in the air sample



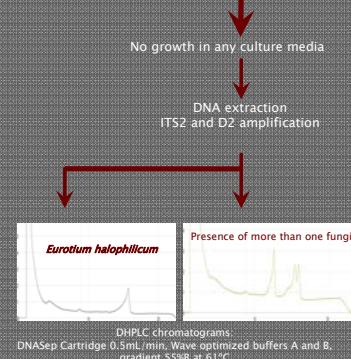
Stachybotrys sp. a potentially toxicogenic fungi producing:



Chaetomium sp. and *Chrysosporium* sp. (400x)



Curvularia sp. and *Trichothecium* sp. (400x)



So far...

1. The chemical assessment performed alerted for values above the national norm (annex VII, decree-law n. 79/2006) regarding human health (O₃ and VOCs). For conservation purposes, the ozone level was also found to be higher than desirable.
2. The biological assessment of the environment yielded the identification of *Stachybotrys* sp. (potentially toxinogenic) which was followed by remedial action. This same assessment in documents surfaces made it possible to identify keratinophytic and cellulolytic fungi.
3. The DHPLC method is a non-labour intensive method for resolving complex mixtures of fungal DNA.

Bibliography

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