

## Exposure to fungi, bacteria, and endotoxin in a museum, with staff reporting work-related symptoms

Anne Mette Madsen, Margit W. Frederiksen & Hjördís Birna Árnadóttir

To cite this article: Anne Mette Madsen, Margit W. Frederiksen & Hjördís Birna Árnadóttir (2025) Exposure to fungi, bacteria, and endotoxin in a museum, with staff reporting work-related symptoms, *Journal of Occupational and Environmental Hygiene*, 22:9, 713-725, DOI: [10.1080/15459624.2025.2499599](https://doi.org/10.1080/15459624.2025.2499599)

To link to this article: <https://doi.org/10.1080/15459624.2025.2499599>



© 2025 The Author(s). Published with  
license by Taylor & Francis Group, LLC



[View supplementary material](#)



Published online: 28 May 2025.



[Submit your article to this journal](#)



Article views: 741



[View related articles](#)



[View Crossmark data](#)

## Exposure to fungi, bacteria, and endotoxin in a museum, with staff reporting work-related symptoms

Anne Mette Madsen , Margit W. Frederiksen , and Hjördís Birna Árnadóttir 

National Research Centre for the Working Environment, Copenhagen Ø, Denmark

### ABSTRACT

Occupational exposure to airborne fungi in indoor environments may pose respiratory health risks. This study aimed to characterize exposure to fungi, bacteria, and endotoxin in a historical museum where the staff reported airway symptoms. Air samples were collected during three separate workdays using personal and stationary samplers. Fungi and bacteria were quantified and identified using MALDI-TOF MS, and the total inflammatory potential was measured through Reactive Oxygen Species (ROS) production in a human cell line. The geometric mean of staff exposure was  $5.9 \times 10^3$  CFU/m<sup>3</sup> fungi,  $1.8 \times 10^3$  CFU/m<sup>3</sup> bacteria, and 7.93 EU/m<sup>3</sup> endotoxin. Staff reported symptoms of the airways, with immediate symptoms upon entering two departments, which exhibited elevated fungal concentrations compared to other departments. The most prevalent fungal species were *Aspergillus conicus*, *A. domesticus*, *A. pseudoglaucus*, *A. pseudogracilis*, and *Cladosporium* spp. Concentrations of bacteria were highest when staff were present and without dominance by any particular species. Staff exposure and stationary samples induced ROS production in a cell line, which correlated with concentrations of fungi, bacteria, and endotoxin. Fungi were detected on the museum artifacts, and concentrations of fungi and bacteria increased during handling of the artifacts. In conclusion, staff in the two departments where airway symptoms were reported were exposed to  $2 \times 10^4$  to  $7 \times 10^4$  CFU/m<sup>3</sup> of fungi, primarily *Aspergillus* spp. and *Cladosporium* spp. This exposure constituted both xerophilic species and species commonly found in water-damaged buildings, and they seemed to have developed on artifacts. These findings underscore the importance of developing preventive strategies to protect the health of museum staff. Moreover, it highlights the challenge of managing fungi adapted to varying humidity conditions.

### KEYWORDS

Maldi-TOF MS; nasal symptoms; ODTs; occupational health; personal exposure; xerophilic fungi

## Introduction

Exposure to fungi in indoor air can constitute a potential risk factor for respiratory problems. In an arts and crafts museum where books were handled, an employee exposed to high concentrations of fungi repeatedly experienced symptoms similar to those of Organic Dust Toxic Syndrome (ODTS) (Kolmodin-Hedman et al. 1986). Other cases include Hypersensitivity Pneumonitis (HP) associated with exposure to the yeast *Aureobasidium pullulans* from an indoor hydroponic system (Engelhart et al. 2009) and with *Fusarium vasicinctum* from walls in a mobile home, which the affected individual was self-renovating (Dickson and Tankersley 2015). In an epidemiological study, 30%

of 103 museum staff members were sensitized to at least one fungal allergen, with prolonged museum employment of more than 5 years identified as a risk factor (Wiszniewska et al. 2009).

Fungi have been documented to colonize various materials within museum artifacts (Kavkler et al. 2015; Nigam et al. 1994; Okpalanozie et al. 2018; Micheluz et al. 2016; Abe 2010; Bastholm et al. 2021; de Carvalho et al. 2018), with airborne fungi detected in museum rooms, repositories, archival storage facilities, and cultural heritage buildings (Rojas and Aira 2012; Bastholm et al. 2022; Ilies et al. 2023; Zielińska-Jankiewicz et al. 2008; Awad et al. 2020; de Carvalho et al. 2018). Cases of fungal presence on museum artifacts have been associated with water damage (Pinheiro et al. 2019;

**CONTACT** Anne Mette Madsen   National Research Centre for the Working Environment, Lersø Parkallé 105, 2100 Copenhagen Ø, Denmark.

 Supplemental data for this article can be accessed online at <https://doi.org/10.1080/15459624.2025.2499599>. AIHA and ACGIH members may also access supplementary material at <http://oeh.tandfonline.com>.

© 2025 The Author(s). Published with license by Taylor & Francis Group, LLC

This is an Open Access article distributed under the terms of the Creative Commons Attribution License (<http://creativecommons.org/licenses/by/4.0/>), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited. The terms on which this article has been published allow the posting of the Accepted Manuscript in a repository by the author(s) or with their consent.

Sterflinger 2010), but also with xerotolerant and xerophilic fungi that can grow with low water activity have been repeatedly found on museum artifacts (Bastholm et al. 2022; Abe 2010; Zalar et al. 2023; Hagiuda et al. 2022) and library books (Micheluz et al. 2015). However, the potential health effects of exposure to these xerophilic fungi have not been investigated.

Few studies have measured airborne bacteria in museum-related environments (Saridaki et al. 2023; Gorny et al. 2016). In occupational settings where large amounts of organic materials are handled, bacterial endotoxin exposure has been associated with symptoms of the airways and ODTs (Basinas et al. 2012). Moreover, endotoxin may have synergistic effects with fungal  $\beta$ -glucan (Engstad et al. 2002). No studies have been found on endotoxin exposure in museum settings; however, air concentrations have been documented previously in residential (Frankel et al. 2012) and educational (Lai et al. 2015) settings.

Exposure to fungi, endotoxin, and bacteria can induce inflammation, with reactive oxygen species (ROS) production playing a pivotal role in the ensuing immune response (Natarajan et al. 2019). ROS serve as key signaling molecules in inflammatory disorders (Mittal et al. 2014), contributing significantly to the pathogenesis of conditions such as asthma, ODTs, and HP by promoting inflammation, tissue damage, and disease progression in the lungs. ROS production upon exposure to airborne microorganisms and dust from occupational settings can be studied *in vitro* in a human promyelocytic leukemia cell line (HL60) (Frankel et al. 2014). This method offers the advantage of evaluating ROS production in response to the complete mixture of airborne particles encountered during a workday. During a workday, employees are typically exposed to several different airborne fungi and bacteria, endotoxin, and other particles (Vitte et al. 2022; Rasmussen et al. 2021).

A recent review describes a lack of quantitative studies linking indoor exposure to fungi directly to harmful health effects (Perez et al. 2024). In this study, the staff exposure in a historical museum setting was characterized. Measurements were conducted as personal exposure samples during work tasks and stationary area samples in various work areas. In addition, surface samples from artifacts were analyzed as potential sources of exposure. An advanced method for identification of fungal and bacterial species, Matrix-Assisted Laser Desorption/Ionization Time-of-Flight Mass Spectrometry (MALDI-TOF-MS), was used in this study. Additionally, endotoxin levels were measured, and the Total Inflammatory Potential (TIP) of staff exposures was assessed using a human

cell line. The aim is to obtain knowledge about occupational bioaerosol exposures in a museum where staff members experienced symptoms in the airways.

## Methods

### *The museum*

The studied museum is a local historical museum dedicated to preserving and communicating cultural heritage. The museum was temporarily located in containers, some of which were accessible to the public while others served as storage repositories. The departments of the museum included in this study were Office 1, where museum artifacts were also present, Office 2, where the staff was often present, and four museum departments with artifacts (Departments 1-4). Each department was located in a separate container. Visible fungi were observed on artifacts in Departments 1 and 2. During the third day of measurement, an air purification system was active with a HEPA filter mounted to reduce particle concentrations in Department 2.

### *Staff and questionnaire*

Contact with the museum was obtained due to a staff member experiencing airway symptoms, suspected to be caused by fungal exposure. The museum staff consisted of a mixture of employees and volunteers (part-time staff). The staff was preparing to move all of the museum's activities to a renovated facility at the time of measurement, thus, the staff mainly worked on cleaning the current facility and artifacts and packing them for moving. Measurements were made during three separate days in late summer 2023. During the first two days of measurements, one person was at work, and during the third day, five people were at work.

Seven staff members participated in the study and were given a questionnaire on skin and airway-related symptoms occurring within the last week and/or 12 months and on the use of protective equipment. The questionnaire was based on previously used questionnaires (Rasmussen et al. 2023).

### *Exposure assessment*

To determine personal exposure, participants were fitted with backpacks containing pumps attached to two personal air samplers (Gesamtstaubprobenahme sampler (GSP), BIG Inc., USA) attached on the front of the backpack, i.e., in the inhalation zone of the participants (14031 E 2021). A 37 mm polycarbonate filter

(pore size 0.8 µm, SKC) was mounted in one sampler, and a 37 mm Teflon filter (pore size 1.0 µm, Merck) was mounted in the other. The sample flow was adjusted twice before sampling and kept at a flow rate of 3.5 L/min. The average sampling time was 169 min.

#### **Stationary samples, outdoor references, temperature, and humidity**

Stationary samples were collected using GSP samplers in four areas during each of the three days: Office 2, Departments 2, 3, and 4. The average sampling time was 333 min.

To determine potential exposures in Department 1 before and during packing of artifacts on day 3 and to get information about the sizes of particles (aero-dynamic diameter) with airborne microorganisms, stationary samples were collected using an Andersen six-stage cascade impactor (ACI; Thermo Fisher Scientific Inc. Waltham, MA, USA) with a flow rate of 28.3 L/min. The ACI sampled bioaerosols into six size fractions. Stage 1 (particle size range for nasal/oral cavity exposure): 7.0 to 12 µm, stage 2 (Pharynx): 4.7 to 7.0 µm, stage 3 (Trachea and primary bronchi): 3.3 to 4.7 µm, stage 4 (secondary bronchi): 2.1 to 3.3 µm, stage 5 (terminal bronchi): 1.1 to 2.2 µm, and stage 6 (Alveoli): 0.65 to 1.1 µm. The ACI was placed on a 1 m stand and sampled for 10 and 20 min on Dichloran glycerol agar (DG18) twice, 15 and 20 min on Malt yeast 50% agar (MY50) twice, and 10 and 15 min on Nutrient agar (NA; Thermo Fisher Scientific Oxoid, Basingstoke, UK) plates with actidione (cycloheximide; 50 mg/L; Serva, Germany) twice. The agar plates from the ACI were incubated at 25 °C upon return to the laboratory.

Outdoor reference samples were taken outside the museum using a GSP sampler mounted with a polycarbonate filter; the sampling time was 330 min. Temperature and relative humidity (RH) were measured inside different departments and outdoors using Tinytag Plus Range (TGP-1500) every 5<sup>th</sup> minute throughout the sampling periods.

#### **Surface samples**

Surface samples were taken from 10 artifacts from different departments using a moistened swab (eSwabs pre-moistened in modified Amies medium; Copan). The artifacts included: paper, a glass cabinet, wooden shelf, Southwest rain hat, typewriter, book, and navigation system.

#### **Extraction, quantification, and identification of bacteria and fungi**

Polycarbonate filters from both personal and stationary GSP samplers were extracted the morning following the sampling day due to the long transport time from the sampling location to the laboratory. Filters were extracted at room temperature in 5.0 mL sterile extraction solution (MilliQ water with 0.05% Tween 80 and 0.85% NaCl) by orbital shaking at 500 rpm for 15 min. Suspensions from the filters were plated in serial dilutions for quantification and identification of bacteria and fungi. For enumeration of bacteria, samples were plated on NA and incubated at 25 °C for seven days. For enumeration of fungi, samples were plated on Dichloran Glycerol agar (DG18; water activity  $a_w = 0.95$ ; Thermo Fisher Scientific Oxoid, Basingstoke, UK) in duplicate and incubated at 25 and 37 °C for seven days. Fungi were also plated on Malt Yeast Extract 50% Glucose Agar (MY50 agar,  $a_w = 0.86$ ; Oxoid, Denmark) in duplicate and incubated at 15 and 25 °C for 14 days.

E-swabs (surface samples) were vortexed for 5 min and plated in 5 serial dilutions on MY50 agar in duplicate, followed by incubation at 15 and 25 °C for 14 days.

All visible bacterial and fungal colonies were counted after the specified incubation period. For GSP and ACI samples, concentrations were calculated as time-weighted averages of colony-forming units (CFU)/m<sup>3</sup>, taking into account the sampling time and the flow rate. Sample concentrations were then calculated as geometric mean (GM) values of appropriate sample dilutions (plates with a CFU count between 1 and 100).

A representative dilution, i.e., with optimal coverage and separation of individual colonies, was used for species identifications using matrix-assisted laser desorption/ionization time-of-flight mass spectrometry (MALDI-TOF MS). A Microflex LT mass spectrometer (Bruker Daltonics) was used using the Bruker Biotyper 3.1 software, with the BDAL library for bacteria (V11) and the Filamentous Fungi library (V4) for fungi. The fungal library was extended with xerophilic *Aspergillus* species. Bacteria were identified using the extended direct transfer method, and fungi were identified using a modified ethanol extraction protocol. The instrument was calibrated weekly using a bacterial test standard (Bruker Daltonics). Isolates were analyzed in duplicates on the MALDI-TOF MS, and the following cutoffs were used for species identifications: isolates with scores of 1.80 or higher were identified to species level, isolates with scores between

1.70 and 1.79 were identified to genus level, and isolates with scores lower than 1.70 were unidentified.

### Total inflammatory potential (TIP) and endotoxin

Suspensions from GSP polycarbonate filters were used in duplicate to determine the total inflammatory potential (TIP) of the samples using a granulocyte-like cell assay (HL-60 cells; ATCC, CCL-240). As negative reference, the extraction solution for polycarbonate filters was used, a positive references, pure endotoxin (0.5 EU/ml of Escherichia coli O55:B5 endotoxin, Lonza Pharma & Biotech), an air sample from a home with more fungi than bacteria, and a sample from a farm with more bacteria than fungi were used. The three positive references were used as internal positive references in every assay.

The HL-60 cells were exposed to the dust suspensions and references, after which the production of reactive oxygen species (ROS) was measured by a luminol-dependent chemiluminometric assay using a thermostatic ( $37^{\circ}\text{C}$ ) ORION II Microplate Luminometer (Berthold Detection Systems) (Madsen et al. 2020, Timm et al. 2006). Relative-light units per second (RLU/sec) were measured for 1 sec every 120 sec for 180 min. Data are presented as ROS production over time, together with concentrations of fungi, bacteria, and endotoxin (units/mL) in the same samples. In addition, for each sample, RLU over the full period of 180 min was summed, thereby expressing the TIP of the sample as the area under the curve (AUC). The AUC was expressed as time-weighted averages ( $\text{AUC}/\text{m}^3$ ).

The dust on the Teflon filters was extracted in 5.0 mL pyrogen-free water with 0.05% Tween 20 by orbital shaking (500 rpm) at room temperature for 60 min and centrifuging ( $1000 \times g$ ) for 15 min. The supernatant was stored at  $-80^{\circ}\text{C}$  until they were diluted 1.5 times and analyzed (in duplicate) for endotoxin using the kinetic Limulus Amebocyte Lysate test (Kinetic-QCL endotoxin kit, Lonza Pharma & Biotech, Walkersville, Maryland, USA). A standard curve obtained from an Escherichia coli O55:B5 reference endotoxin was used to determine the concentrations in terms of endotoxin units (EU;  $11.0 \text{ EU} \approx 1.0 \text{ ng}$ ). The data are presented as  $\text{EU}/\text{m}^3$  air, and the limit of detection was 0.05 EU/ml, corresponding to a detection limit of  $0.085 \text{ EU}/\text{m}^3$ .

### Data analyses

Data were  $\log_{10}$ -transformed before analysis to approximate normality and stabilize variance, as raw

concentration data were right-skewed. After transformation, the data were visually assessed for normality using Q-Q plots, and residuals from the General Linear Model (GLM) were checked to confirm that model assumptions (normality and homogeneity of variance) were met. To compare exposures and concentrations measured using a stationary sampler in different areas of the museum, the GLM procedure in SAS was used. The LSD (Least Significant Difference) option was used to conduct pairwise comparisons of means. Concentrations before and during activity with packing of artifacts, and the geometric mean aerodynamic diameter of particles with bacteria on NA and fungi on DG18 and My50 were compared using the paired t-test. Results were considered statistically significant at  $p < 0.05$ . Pearson correlations were calculated to assess the relationship between exposure and TIP. Statistical analyses were done using SAS 9.4 (Cary, North Carolina, USA).

## Results

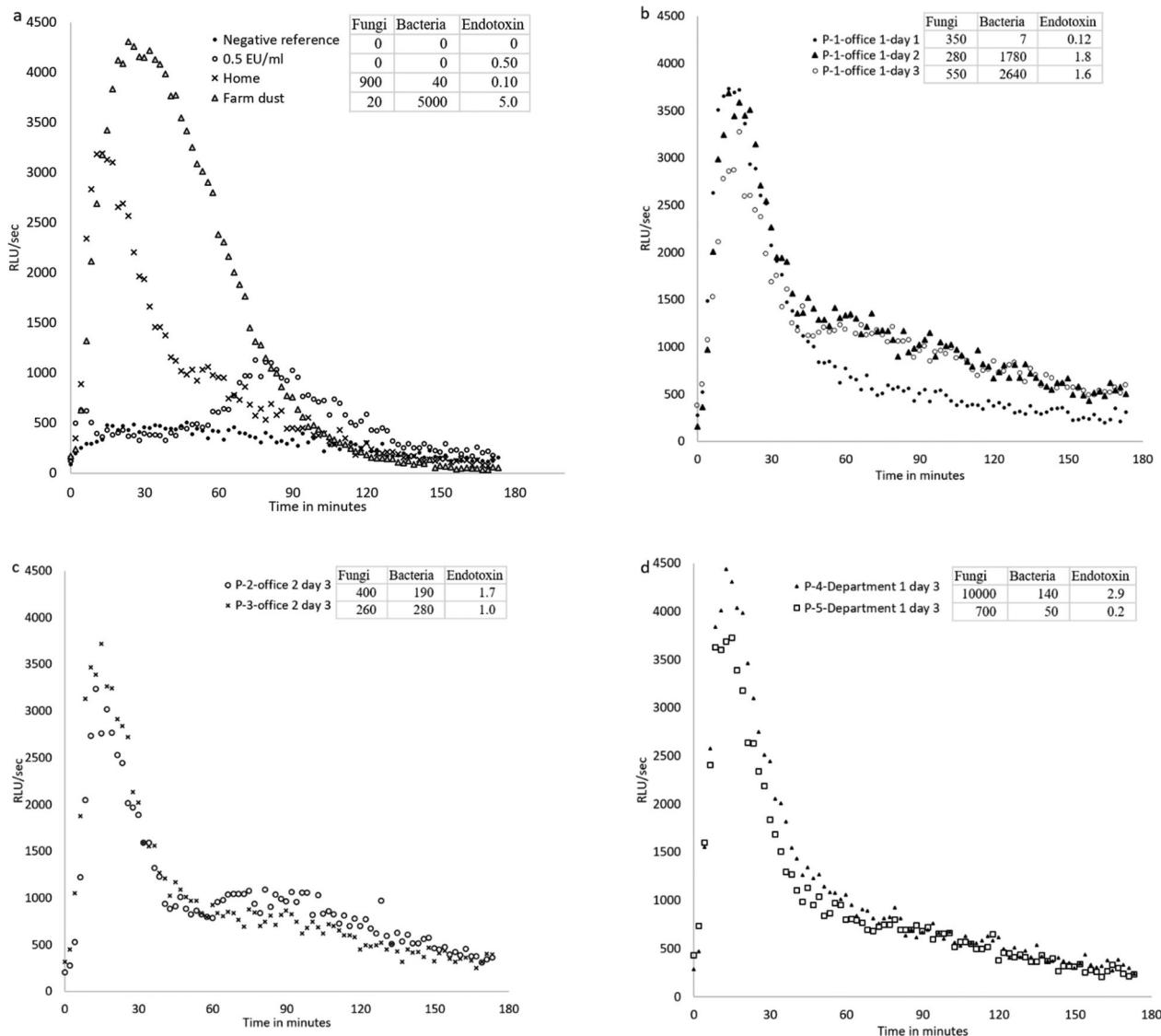
### **Self-reported symptoms**

The staff were nonsmokers, and more than 50% reported nasal symptoms that resolved after leaving work. Some reported sneezing at work and the sudden onset and resolution of influenza-like symptoms (Table S1). The staff reported immediate airway symptoms when entering Departments 1 and 2. While staff often used gloves at work, facial masks were used only after symptoms appeared.

### **TIP, exposure, and concentrations of microorganisms**

The average RH was higher than 60% in all departments, and indoor temperatures were affected by outdoor temperatures (Table S2).

Profiles of ROS production of HL60 cells during exposure to personal and stationary samplers are presented in Figures 1 and 2, respectively, without normalization to sampling time. Negative (extraction solution) and positive reference samples—endotoxin, a sample from a home, and a sample from a farm are shown in Figure 1. All museum samples induced ROS production. No peak response was observed for the negative reference, while peak responses were observed after 75 min for pure endotoxin, 13 min. for the home sample, and 23 min for the farm sample. Peak responses were observed 10–15 min. after exposure to the personal (Figure 1b–d) and stationary (Figure 2a–d) museum samples. In samples with the highest



**Figure 1.** Profiles of reactive oxygen species (ROS) production in HL60 cells during exposure of cells to reference and personal air samples. The response is measured in relative luminescence units (RLU) per second over time. Each panel includes a table displaying concentrations of fungi (cfu/ml), bacteria (cfu/ml), and endotoxin (EU/ml) in the corresponding sample. Shown are results for: one person working in office 1 over three days (b); two persons in office 2 (c); and two persons in department 1 (d).

concentrations of bacteria or endotoxin, ROS production persisted beyond 60 min (Figure 1b). The highest peak response was found for P-4, who worked in Department 3 for 178 min. (Figure 1d). For the stationary samples, the highest peak exposure was observed in Department 2 on days 1 and 2 (Figure 2b).

The GM of TIP, fungi, bacteria, and endotoxin in personal exposures were  $8.20 \times 10^5$  AUC/m<sup>3</sup>,  $5.9 \times 10^3$  CFU/m<sup>3</sup>,  $1.8 \times 10^3$  CFU/m<sup>3</sup>, and 7.93 EU/m<sup>3</sup>, respectively. The TIP ( $p < 0.01$ ), bacteria ( $p < 0.01$ ), and endotoxin ( $p < 0.01$ ) of personal exposures were higher than in stationary samples.

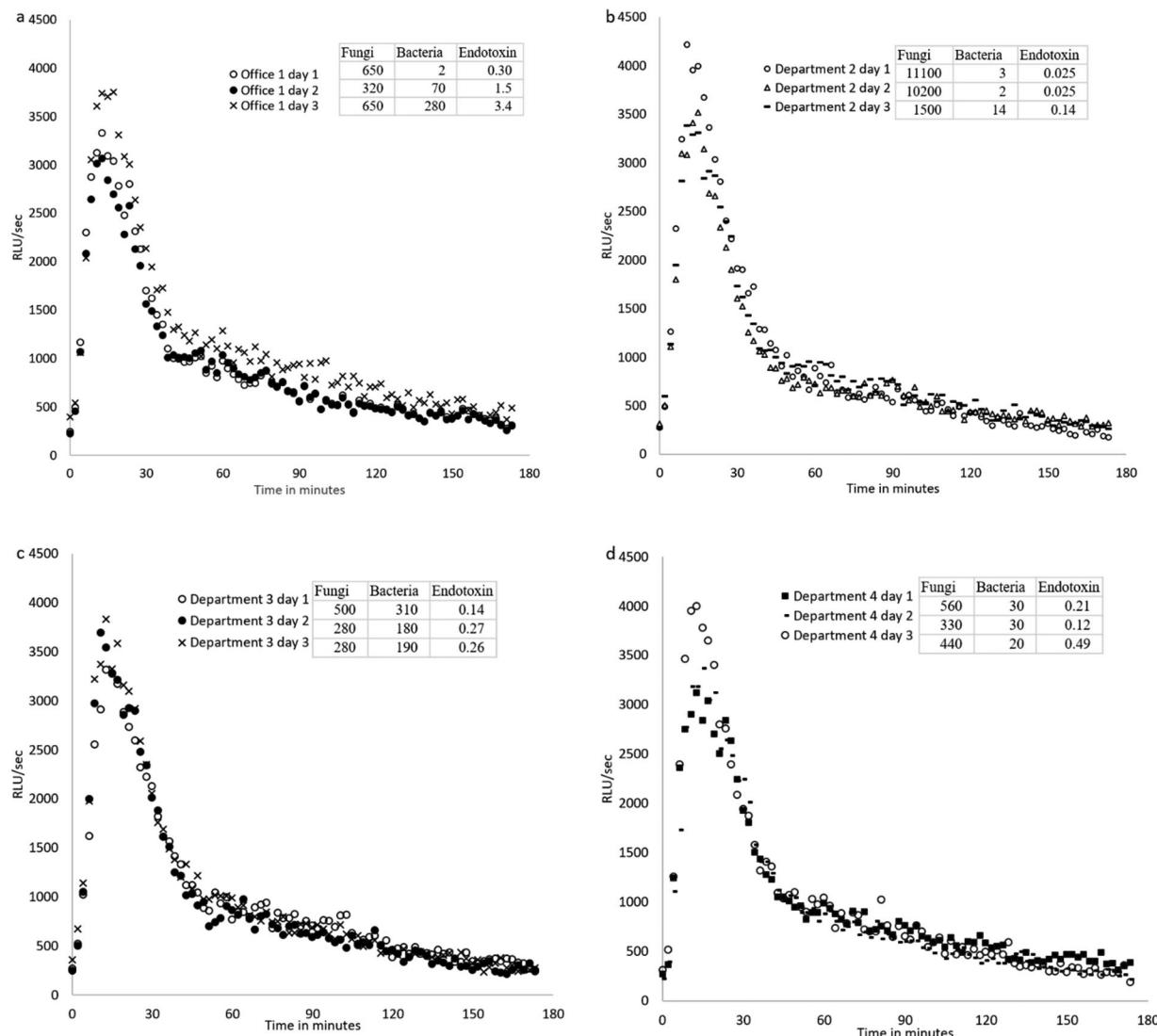
The staff were exposed to different levels of fungi, bacteria, and endotoxin when working in different

departments, with the highest TIP, fungal, and endotoxin exposure when working in Department 1, and the highest bacterial exposure in Office 1. High concentrations of fungi were found in Department 2 (Table 1).

TIP correlated with endotoxin ( $r = 0.72$ ,  $p < 0.01$ ), fungi ( $r = 0.44$ ,  $p = 0.05$ ), and bacteria ( $r = 0.87$ ,  $p < 0.01$ ). Endotoxin correlated with concentrations of bacteria ( $r = 0.84$ ,  $p < 0.01$ ).

#### Fungal and bacterial species

Several *Aspergillus* and *Penicillium* species were found in the air samples, and some are described as



**Figure 2.** Profiles of reactive oxygen species (ROS) production in HL60 cells during exposure of cells to air samples taken in office 1 (a), departments 2, 3, and 4 (b, c, and d) using stationary samplers. The response is measured in relative luminescence units (RLU) per second over time. Each panel includes a table displaying concentrations of fungi (cfu/ml), bacteria (cfu/ml), and endotoxin (EU/ml) in the corresponding sample.

xerophilic (Table 2). Exposure during work in Department 1 was associated with exposures to *Aspergillus conicus*, *A. pseudoglaucus*, and *A. domesticus*. In all areas, airborne *Cladosporium* was found (Table 2).

The fungal species: *A. conicus*, *A. niger*, *Cladosporium* sp., *P. camemberti*, *P. chrysogenum*, *P. commune*, *P. corylophilum*, *Penicillium* sp., and *Saccharomyces cerevisiae* (baker's yeast) were found on the artifacts.

The staff was exposed to different species of bacteria, including human skin-related species such as *Micrococcus luteus*, *Moraxella osloensis*, *Staphylococcus epidermidis*, and *S. hominis*, which were also found using stationary samplers in Office 1 (Table 3).

### Packing of artifacts

Using the ACI in Department 1, airborne fungi and bacteria were sampled in six size fractions before and during work with packing artifacts (Tables S3 and S4; Figure 3). Concentrations of fungi and bacteria were higher during the activity than before ( $p < 0.05$ ), and the fungal concentration increased by a factor of 6.6 while bacteria increased by a factor of 2.8. Several of the fungal species found during the packing activity were also found before, but in lower concentrations. The geometric mean aerodynamic diameter of fungal spores before the packing activity was  $2.6 \mu\text{m}$  and during the activity,  $2.8 \mu\text{m}$  ( $p > 0.05$ ). Before the work activity, the geometric mean aerodynamic diameter of

particles with bacteria was  $2.8 \mu\text{m}$ , and during the activity,  $3.3 \mu\text{m}$  ( $p < 0.05$ ).

## Discussion

In this study, bioaerosol exposure of part-time staff in a historical museum was characterized. Staff members

**Table 1.** TIP, exposure and concentrations (GM) inside and outside the museum; number in the same column followed by the same letter are not statistically different.

|                                      | TIP<br>AUC/m <sup>3</sup> | Fungi<br>CFU/m <sup>3</sup> | Bacteria<br>CFU/m <sup>3</sup> | Endotoxin<br>EU/m <sup>3</sup> |
|--------------------------------------|---------------------------|-----------------------------|--------------------------------|--------------------------------|
| P-value                              | <0.01                     | <0.01                       | 0.07                           | <0.01                          |
| Office 1 P (n = 3)                   | $3.54 \times 10^{5b}$     | 2665 <sup>b</sup>           | 2217 <sup>a</sup>              | 4.8 <sup>a</sup>               |
| Office 2 P (n = 2)                   | $3.46 \times 10^{5b}$     | 2804 <sup>b</sup>           | 1980 <sup>a</sup>              | 11.6 <sup>a</sup>              |
| Department 1 <sup>1)</sup> P (n = 2) | $6.18 \times 10^{5a}$     | $4.13 \times 10^{4a}$       | 1289 <sup>a</sup>              | 11.4 <sup>a</sup>              |
| Office 1 S (n = 3)                   | $1.91 \times 10^{5c}$     | 2098 <sup>b</sup>           | 129 <sup>ab</sup>              | 5.4 <sup>a</sup>               |
| Department 2 <sup>2)</sup> S (n = 3) | $1.03 \times 10^{5c}$     | $2.30 \times 10^{4a}$       | 18 <sup>b</sup>                | 0.18 <sup>bc</sup>             |
| Department 3 <sup>3)</sup> S (n = 3) | $1.15 \times 10^{5c}$     | 1416 <sup>b</sup>           | 31 <sup>b</sup>                | 0.86 <sup>b</sup>              |
| Department 4 <sup>4)</sup> S (n = 3) | $1.28 \times 10^{5c}$     | 1774 <sup>b</sup>           | 113 <sup>b</sup>               | 0.95 <sup>b</sup>              |
| Outdoor S (n = 3)                    | $5.02 \times 10^{4d}$     | 316 <sup>c</sup>            | 208 <sup>b</sup>               | 0.19 <sup>c</sup>              |

P = personal samples, S = stationary samples, n = numbers of samples.<sup>1)</sup> Visible fungi on the artifacts; human activity.<sup>2)</sup> Most artifacts in glass display cases; visible fungi on artifacts; an air purification system was active during day 3, no human activity in the department.<sup>3)</sup> Mainly wooden bookcases with mugs; no visible fungi; no human activity.<sup>4)</sup> Styled like a formal historical living room, furnished with vintage wood and fabric furniture; no visible fungi; no human activity.

reported nasal symptoms and symptoms resembling those of ODTs. Elevated exposure to fungi was linked to immediate airway symptoms upon entering specific departments. Exposure levels to fungi (maximum in Department 1:  $7 \times 10^4 \text{ CFU/m}^3$ ) were lower than those reported in a case of museum-related ODTs ( $10^6 \text{ CFU/m}^3$ ) (Kolmodin-Hedman et al. 1986). In this study, the fungal species *Aspergillus conicus*, *A. domesticus*, *A. pseudoglaucus*, *A. pseudogracilis*, and *Cladosporium* spp. dominated the exposures and may be the primary causative agents of the reported symptoms, and exposure to endotoxin and bacteria may also contribute. Thus, ROS, which are key signaling molecules in inflammatory disorders, correlated with airborne concentrations of fungi, bacteria, and endotoxin. High concentrations of fungal spores were associated with the highest peak ROS production, as seen for the exposure of P4 working in Department 1, while the highest bacterial and endotoxin concentrations seemed to be associated with a prolonged ROS production, as seen for P1 during working in office 1. Previous studies have shown that exposure to endotoxin may act synergistically with fungal exposure to exacerbate respiratory inflammation (Park et al. 2006).

**Table 2.** Occupational exposure<sup>1)</sup> to fungal species during work in offices 1 and 2 and department 1. Stationary measurements in office 2 and other departments of the museum as well as outdoors. Data presented as averages of positive samples in CFU/m<sup>3</sup>.

| Fungal species  | Office 1<br>P (n = 3) | Office 2<br>P (n = 3) | Depart 1<br>P (n = 2) | Office 1<br>S (n = 3) | Depart. 2<br>S (n = 3) | Depart 3<br>S (n = 3) | Depart 4<br>S (n = 3) | Positive<br>Samples | Out-door<br>References |
|---|-----------------------|-----------------------|-----------------------|-----------------------|------------------------|-----------------------|-----------------------|---------------------|------------------------|
| <i>Alternaria chartarum</i> <sup>x</sup>                          |                       |                       |                       |                       |                        | 45                    | 44                    | 3                   | 3                      |
| <i>Al. infectoria</i> <sup>x</sup>                                |                       |                       |                       |                       |                        |                       |                       | 1                   |                        |
| <i>Aspergillus conicus</i> <sup>H,X</sup>                         | 51                    |                       |                       | 23                    |                        |                       |                       | 5                   |                        |
| <i>A. creber</i>  |                       |                       |                       | 44                    | 4700                   | 45                    |                       |                     |                        |
| <i>A. domesticus</i> <sup>x</sup>                                 |                       |                       |                       | 400                   |                        |                       |                       | 1                   |                        |
| <i>A. fumigatus</i> <sup>rc2, H</sup>                             |                       |                       |                       | 5058                  | 5600                   |                       |                       | 3                   |                        |
| <i>A. glaucus</i> <sup>x</sup>                                    |                       |                       |                       |                       | 28                     |                       |                       | 1                   |                        |
| <i>A. montevidensis</i> <sup>x</sup>                              | 94                    |                       |                       |                       |                        | 76                    |                       | 1                   |                        |
| <i>A. niger</i>   | 25                    |                       |                       |                       |                        |                       |                       | 1                   |                        |
| <i>A. pseudoglaucus</i> <sup>x</sup>                              | 51                    |                       |                       |                       |                        |                       |                       | 1                   |                        |
| <i>A. pseudogracilis</i> <sup>x</sup>                             | 25                    |                       |                       | 4289                  | 2022                   | 44                    |                       | 7                   |                        |
| <i>A. versicolor</i> <sup>rc2, H</sup>                            |                       |                       |                       | 4421                  | 2400                   |                       |                       | 3                   |                        |
| <i>A. villosus</i> <sup>x</sup>                                   |                       |                       |                       |                       | 103                    | 66                    |                       | 4                   |                        |
| <i>Botrytis cinerea</i>   |                       |                       |                       |                       | 103                    |                       |                       | 1                   |                        |
| <i>Cladosporium cladosporioides</i> <sup>x</sup>                  |                       | 91                    |                       |                       |                        |                       |                       | 3                   |                        |
| <i>C. cucumerinum</i>   |                       |                       |                       |                       |                        | 79                    |                       | 3                   |                        |
| <i>C. herbarum</i> <sup>x</sup>                                   | 91                    |                       |                       |                       |                        |                       |                       | 26                  |                        |
| <i>Cladosporium</i> sp. <sup>x</sup>                              | 61                    | 151                   |                       | 26                    | 39                     | 32                    |                       | 8                   |                        |
| <i>C. sphaerospermum</i> <sup>x</sup>                             | 900                   | 349                   | 810                   | 617                   | 789                    | 309                   | 469                   | 17                  |                        |
| <i>Penicillium brevicompactum</i> <sup>H,X</sup> / <i>olsonii</i> | 300                   |                       |                       |                       |                        |                       |                       | 1                   |                        |
| <i>P. camemberti</i> <sup>H</sup>                                 | 128                   | 91                    |                       | 210                   | 76                     | 46                    | 85                    | 9                   | 3                      |
| <i>P. chrysogenum</i> <sup>X</sup>                                | 970                   |                       |                       |                       |                        |                       |                       | 2                   |                        |
| <i>P. citreonigrum</i> <sup>X</sup>                               | 158                   |                       | 4221                  | 5                     |                        |                       | 87                    | 6                   |                        |
| <i>P. citrinum</i>  |                       |                       |                       |                       |                        | 15                    |                       | 3                   |                        |
| <i>P. commune</i> <sup>H</sup>                                    | 5                     |                       | 190                   | 23                    | 152                    | 92                    |                       | 8                   |                        |
| <i>P. corylophilum</i> <sup>H,X</sup>                             | 180                   | 78                    |                       | 65                    |                        | 227                   |                       | 6                   | 26                     |
| <i>P. digitatum</i>   |                       |                       |                       |                       |                        |                       |                       | 1                   |                        |
| <i>Penicillium</i> sp.  |                       |                       |                       | 4221                  |                        | 5320                  |                       | 4                   |                        |
| <i>Wallemia sebi</i> <sup>H,X</sup>                               |                       |                       |                       |                       |                        | 170                   |                       | 1                   |                        |

<sup>1)</sup>P = personal samples, S = stationary samples, n = numbers of samples. <sup>x</sup>Fungi considered as xerophilic or xerotolerant, <sup>H</sup>Described to be associated with moisture damage buildings (Vesper et al. 2007), <sup>rc2</sup>Risk class 2 pathogen. See also footnotes in Table 1. Positive samples are numbers of positive samples in the museum. Concentrations are presented as a heatmap with darker red presenting highest concentrations.

**Table 3.** Occupational exposure<sup>1)</sup> to bacterial species (CFU/m<sup>3</sup>; average of positive samples) during work in offices 1 and 2 and department 2. Stationary measurements in other departments of the museum as well as outdoors.

| Bacterial species                      | Office 1<br>P (n = 3) | Office 2<br>P (n = 3) | Depart1<br>P (n = 2) | Office 1<br>S (n = 3) | Depart 2<br>S (n = 3) | Depart 3<br>S (n = 3) | Depart 4<br>S (n = 3) | Positive<br>Samples | Out-door<br>References |
|--|-----------------------|-----------------------|----------------------|-----------------------|-----------------------|-----------------------|-----------------------|---------------------|------------------------|
| <i>Acinetobacter lwoffii</i>           | 879                   |                       |                      |                       |                       |                       |                       | 1                   |                        |
| <i>Agromyces cerinus</i>               |                       | 38                    |                      |                       |                       |                       |                       | 1                   |                        |
| <i>Arthrobacter luteolus</i>           |                       | 38                    |                      | 15                    |                       |                       |                       | 2                   |                        |
| <i>Bacillus cereus</i>                 |                       | 38                    |                      | 15                    |                       |                       |                       | 2                   |                        |
| <i>B. mycoides</i>                     |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>Brevibacterium casei</i>            |                       | 38                    |                      |                       |                       |                       |                       | 1                   |                        |
| <i>Corynebacterium lipophiloflavum</i> |                       | 79                    |                      |                       |                       |                       |                       | 1                   |                        |
| <i>Co. mucifaciens</i>                 | 16                    | 114                   | 16                   |                       |                       |                       |                       | 5                   |                        |
| <i>Co. ureicelerivorans</i>            |                       | 114                   |                      |                       |                       |                       |                       | 1                   |                        |
| <i>Curtobacterium flaccumfaciens</i>   | 16                    | 38                    |                      | 8                     |                       |                       | 29                    | 5                   |                        |
| <i>Cytobacillus firmus</i>             |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>Exiguobacterium mexicanum</i>       |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>Frigoribacterium faeni</i>          | 16                    |                       | 16                   | 15                    |                       |                       |                       | 3                   |                        |
| <i>Kocuria marina</i>                  |                       |                       |                      | 40                    |                       |                       |                       | 1                   |                        |
| <i>K. palustris</i>                    |                       |                       |                      | 40                    |                       |                       | 15                    | 2                   |                        |
| <i>K. rhizophila</i>                   |                       |                       |                      | 60                    |                       |                       |                       | 1                   |                        |
| <i>K. rosea</i>                        |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>Kytococcus sedentarius</i>          |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>Microbacterium foliorum</i>         |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>M. phyllophagae</i>                 |                       |                       |                      |                       |                       | 11                    |                       | 1                   |                        |
| <i>Micrococcus luteus</i>              | 1758                  | 197                   | 128                  | 450                   | 5                     |                       | 171                   | 10                  |                        |
| <i>Moraxella osloensis</i>             | 1758                  | 310                   | 401                  | 15                    |                       |                       |                       | 6                   |                        |
| <i>Oerskovia turbata</i>               |                       |                       |                      | 5                     |                       |                       |                       | 1                   |                        |
| <i>Paenarthrobacter aurescens</i>      |                       |                       |                      | 16                    |                       |                       |                       | 1                   |                        |
| <i>Planococcus okeanokoites</i>        |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>Pseudomonas oryzihabitans</i>       |                       |                       | 38                   | 40                    |                       |                       |                       | 2                   |                        |
| <i>Rhodococcus erythropolis</i>        |                       |                       |                      | 16                    |                       |                       |                       | 1                   |                        |
| <i>R. fascians</i>                     |                       | 189                   | 64                   | 40                    |                       |                       | 15                    | 6                   | 77                     |
| <i>Staphylococcus arlettae</i>         |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>S. epidermidis</i>                  | 879                   | 159                   | 16                   | 40                    |                       |                       |                       | 6                   |                        |
| <i>S. equorum</i>                      |                       | 76                    |                      |                       |                       |                       |                       | 1                   |                        |
| <i>S. haemolyticus</i>                 |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>S. hominis</i>                      | 5275                  | 250                   | 1001                 | 114                   |                       |                       | 15                    | 8                   |                        |
| <i>S. lugdunensis</i>                  | 253                   |                       |                      |                       |                       |                       |                       | 1                   |                        |
| <i>S. petrasii</i>                     |                       |                       |                      | 8                     |                       |                       |                       | 1                   |                        |
| <i>S. saprophyticus</i>                |                       |                       |                      | 30                    |                       |                       |                       | 1                   |                        |
| <i>S. warneri</i>                      |                       | 79                    |                      |                       |                       |                       |                       | 1                   |                        |
| <i>Streptomyces chartreusis</i>        |                       |                       | 16                   |                       |                       |                       |                       | 1                   |                        |
| <i>S. xanthophaeus</i>                 |                       | 38                    |                      |                       |                       |                       |                       | 1                   |                        |

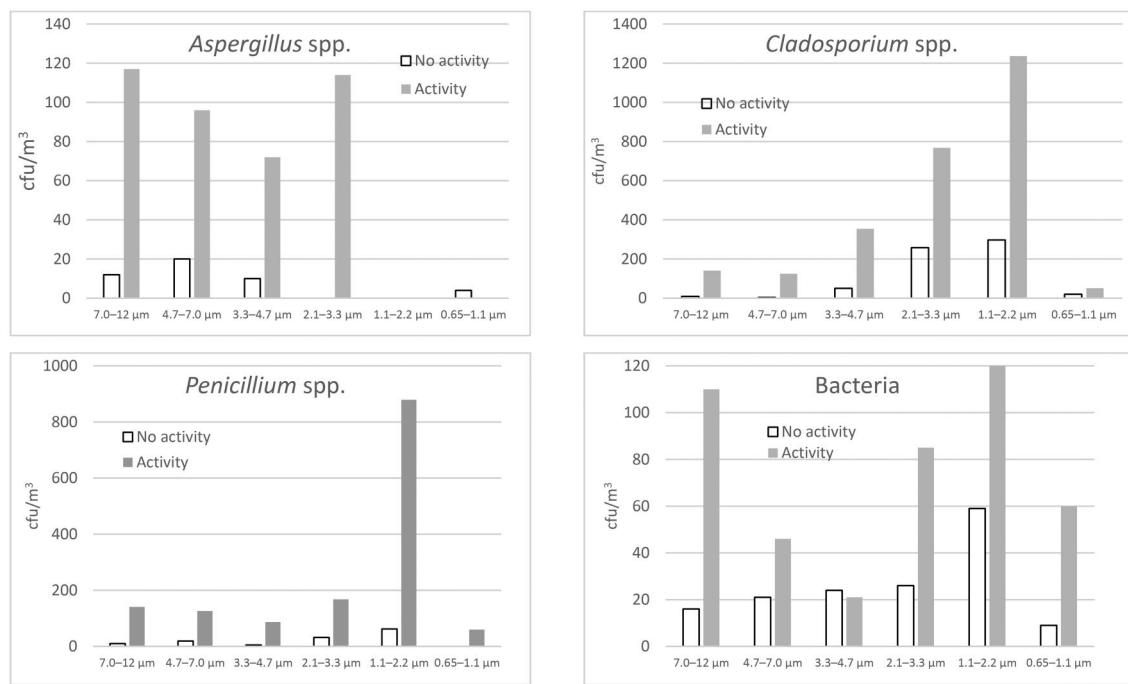
<sup>1)</sup>P = personal and S = stationary samples taken using GSP samplers. See also footnotes in Table 1. Positive samples are numbers of positive samples in the museum. Concentrations are presented as a heatmap with darker red presenting highest concentrations.

High concentrations of airborne *A. conicus* were found in different departments. Rare cases of eye disease in immunocompromised patients have been reported from exposure to this species (Smith et al. 2013; Jones 1978). The species *A. montevidensis*, which was found in low concentrations, has previously been found in clinical airway and clinical sputum samples, and it has been involved in cases of otitis, keratitis, and pulmonary infection (Siqueira et al. 2018; Bongomin et al. 2018). Exposure to *Aspergillus fumigatus* and *A. niger*, found in low concentrations, have previously been shown to cause different occupational related health effects in settings with exposures to greater concentrations than found in this study (Kofoed et al. 2024).

*Aspergillus* spp. was present as airborne particles of the size mainly depositing in the nose, throat, and bronchial region, while *Cladosporium* spp. was present as smaller particles able to deposit in both upper and

lower airways. Hence, they may affect the airways differently.

No other publications were identified on personal exposures to fungi, bacteria, and endotoxins in museum environments. Consequently, the findings are compared with studies using stationary samplers. The fungal concentrations in all departments exceeded or were in the upper range of what has been measured in archive storage facilities and museums (Zielinska-Jankiewicz et al. 2008; Awad et al. 2020; Saridaki et al. 2023; Guiamet et al. 2011), and exposure to endotoxin was higher than measured in normal homes (Frankel et al. 2012), a laboratory (Niu et al. 2020), and outdoor air (Madsen 2006). Personal exposure to bacteria was higher than found in museums in Poland and Greece (Lenart-Boron et al. 2016; Saridaki et al. 2023). In contrast, stationary measurements were lower than reported in these studies. It should be noted that measured concentrations may be affected by the



**Figure 3.** Concentrations of *Aspergillus* spp., *cladosporium* spp., and *penicillium* spp. and bacteria in department 1 as function of particle size fraction. Samples were taken using the ACI before and during work activities with packing of artifacts. Fungal and bacterial species can be found in Tables S3 and S4.

sampling devices and agar media used. For example, some of the cited studies used passive samplers with sedimentation on agar media, whereas this study used the GSP sampler, which actively samples the inhalable fraction of airborne particles (Kenny et al. 1999).

The temperature in the museum was at the lower end of the recommended range (16 to 25 °C) for exhibitions, while the average RH exceeded the recommended range of 40 to 60% (Museernes-Samråd 2021) with some fluctuations in both parameters. These conditions likely exposed artifacts to moisture, facilitating fungal growth. Consistent with this, several of the identified fungal species found in the air and on artifacts are known to be associated with moisture in buildings. However, the fungi found in high concentrations, *A. conicus*, *A. domesticus*, *A. pseudoglaucus*, *Cladosporium cladosporioides*, and *P. chrysogenum* are xerophilic fungi that can grow in environments with low water activity. Interestingly, of these xerophilic species, some are commonly (Vesper et al. 2007; Chakravarty 2022) or occasionally (Andersson et al. 2022) found in humid buildings. In the mentioned case study with ODTs, two species, *A. versicolor* and *P. verrucosum*, were identified (Kolmodin-Hedman et al. 1986). While *A. versicolor* is commonly found in humid buildings, *P. verrucosum* is xerophilic.

Based on the presence of xerophilic species in both wet and dry buildings, strategies to avoid fungal growth are challenged. It has been suggested that xerophilic species may colonize dry material, and later, the dead hyphae of the xerophilic species form the basis for growth of non-xerophilic species (Micheluz et al. 2015). Recommendations to avoid fungal growth in museum artifacts have been suggested to include using fans to foster air movement and controlling temperature and humidity (Koul and Upadhyay 2018).

Some of the species found in the air in this study (*A. niger*, *P. funiculosum*, and *P. corylophilum*) have previously been found in museums and archive storage facilities (Zielińska-Jankiewicz et al. 2008; Bastholm et al. 2024), in indoor air in museums (*A. flavus*, *A. niger*, *C. cladosporioides*, *C. sphaerospermum*, *P. citrinum*, and *P. corylophilum*) (Rojas and Aira 2012), and on wood in a historical building (*A. conicus*) (Slimen et al. 2020). In this study, *Alternaria* spp. was found in low concentrations, but in a historical museum hall, it was found as the dominating genus (Saridaki et al. 2023). Although *Cladosporium* spp. were found both inside and outside the museum departments, higher concentrations were found inside, and they increased during packing activities. This suggests that the airborne *Cladosporium*

spp. concentrations derived from the artifacts, but whether it has colonized the artifacts or was present in sedimented dust is not known.

Several of the bacterial species found in this study are described as plant, soil, and human-related species. Findings of bacteria in the highest concentrations in personal samples, followed by office samples, indicate that bacteria derive from the staff and their activities. This was further supported by the presence of skin-related bacterial species such as *M. luteus* and *Staphylococcus hominis*. In Department 1, concentrations of airborne bacteria increased during activities. In a museum in Italy, concentrations of airborne bacteria increased during opening hours (Pasquarella et al. 2011), and in a study with stationary measurements in different rooms of a historical museum, the highest concentration of airborne bacteria was also found in an office (Grabek-Lejko et al. 2017).

## Limitations

In this study, a combination of personal and stationary samplers was used to obtain knowledge about exposure and concentrations in museum departments with staff reporting work-related symptoms of the airway. A limitation of the study is the low number of participants, though all staff members at work participated in the study. Exposure levels may not reflect concentrations over time. This limitation was difficult to mitigate as the museum was undergoing deinstallation due to fungal contamination. Nevertheless, this study provides important information regarding indoor bioaerosol exposure in a museum facility.

In this study, a combination of three agar media and MALDI-TOF MS was used for the identification of fungi and bacteria. The use of additional agar media may result in the identification of more species.

## Conclusion

The museum staff reporting airway-related symptoms were exposed to elevated levels of fungi and endotoxin. In the two departments where staff reported immediate symptoms, fungal concentrations ranged from 2 to  $7 \times 10^4$  CFU/m<sup>3</sup>. The predominant species were *Aspergillus conicus*, *A. domesticus*, *A. pseudoglaucus*, and *A. pseudogracilis*. Little is known about exposure to these species, and further research is needed to elucidate the health effects of exposure to elevated concentrations.

## Recommendations

The exposure of affected staff consisted of both xerophilic species and those commonly found in water-damaged buildings, highlighting the challenge of managing fungi adapted to varying humidity conditions. This study underscores the importance of developing targeted interventions and preventive strategies to mitigate exposure and protect the health of museum staff. Fungi were found on museum artifacts, and handling these artifacts was associated with increased air concentration of fungi. This suggests that the artifacts serve as a source of exposure, highlighting the importance of limiting fungal growth to protect both staff and artifacts.

## Acknowledgements

We would like to thank all the participants in the study and the funding sources.

## Disclosure statement

No potential conflict of interest was reported by the author(s).

## Funding

This work was supported by Augustinus foundation-22-1543, Aase og Ejnar Danielsens fond-22-30-0478.

## ORCID

Anne Mette Madsen  <http://orcid.org/0000-0002-5895-7521>

Margit W. Frederiksen  <http://orcid.org/0000-0001-7184-1171>

Hjörðís Birna Árnadóttir  <http://orcid.org/0009-0001-5271-6284>

## Data availability statement

Exposure data can be found in the supplementary file.

## References

- 14031 E. 2021. EUROPEAN STANDARD Workplace exposure - Quantitative measurement of airborne endotoxins. EUROPEAN COMMITTEE FOR STANDARDIZATION.
- Abe K. 2010. Assessment of the environmental conditions in a museum storehouse by use of a fungal index. Int Biodeterior Biodegrad. 64(1):32–40. doi: [10.1016/j.ibiod.2009.10.004](https://doi.org/10.1016/j.ibiod.2009.10.004).
- Andersson M, Varga A, Mikkola R, Vornanen-Winqvist C, Salo J, Kredics L, Kocsbá S, Salonen H. 2022. Aspergillus was the dominant genus found during diversity tracking of

- potentially pathogenic indoor fungal isolates. *Pathogens*. 11(10):1171. doi: [10.3390/pathogens11101171](https://doi.org/10.3390/pathogens11101171).
- Awad AHA, Saeed Y, Shakour AA, Abdellatif NM, Ibrahim YH, Elghanam M, Elwakeel F. 2020. Indoor air fungal pollution of a historical museum, Egypt: a case study. *Aerobiologia*. 36(2):197–209. doi: [10.1007/s10453-019-09623-w](https://doi.org/10.1007/s10453-019-09623-w).
- Basinas I, Schlünssen V, Heederik D, Sigsgaard T, Smit LA, Samadi S, Omland O, Hjort C, Madsen AM, Skov S, et al. 2012. Sensitisation to common allergens and respiratory symptoms in endotoxin exposed workers: a pooled analysis. *Occup Environ Med*. 69(2):99–106. doi: [10.1136/oem.2011.065169](https://doi.org/10.1136/oem.2011.065169).
- Bastholm CJ, Andersen B, Frisvad JC, Oestergaard SK, Nielsen JL, Madsen AM, Richter J. 2024. A novel contaminant in museums? A cross-sectional study on xerophilic Aspergillus growth in climate-controlled repositories. *Sci Total Environ*. 944:173880.
- Bastholm CJ, Madsen AM, Andersen B, Frisvad JC, Richter J. 2022. The mysterious mould outbreak—A comprehensive fungal colonisation in a climate-controlled museum repository challenges the environmental guidelines for heritage collections. *J Cult Heritage*. 55:78–87. doi: [10.1016/j.culher.2022.02.009](https://doi.org/10.1016/j.culher.2022.02.009).
- Bastholm CJ, Madsen AM, Frisvad JC, Richter J. 2021. Xerophilic fungi in museum repositories challenge our perception of healthy buildings and the preservation of cultural heritage. *Healthy Buildings 2021–Europe. Proceedings of the 17th International Healthy Buildings Conference* 21–23 June 2021.
- Bongomin F, Moore CB, Masania R, Rowbotham E, Alastrauey-Izquierdo A, Novak-Frazer L, Richardson MD. 2018. Sequence analysis of isolates of Aspergillus from patients with chronic and allergic aspergillosis reveals a spectrum of cryptic species. *Future Microbiol*. 13(14): 1557–1563. doi: [10.2217/fmb-2018-0178](https://doi.org/10.2217/fmb-2018-0178).
- Chakravarty P. 2022. Mycobiota and mycotoxin-producing fungi in southern California: their colonisation and in vitro interactions. *Mycology*. 13(4):293–304. doi: [10.1080/21501203.2022.2104950](https://doi.org/10.1080/21501203.2022.2104950).
- de Carvalho HP, Mesquita N, Trovão J, Rodríguez SF, Pinheiro AC, Gomes V, Alcoforado A, Gil F, Portugal A. 2018. Fungal contamination of paintings and wooden sculptures inside the storage room of a museum: are current norms and reference values adequate? *J Cult Heritage*. 34:268–276. doi: [10.1016/j.culher.2018.05.001](https://doi.org/10.1016/j.culher.2018.05.001).
- Dickson SD, Tankersley MS. 2015. Fatal hypersensitivity pneumonitis from exposure to *Fusarium vasicinfectum* in a home environment: a case report. *Int Arch Allergy Immunol*. 166(2):150–153. doi: [10.1159/000377631](https://doi.org/10.1159/000377631).
- Engelhart S, Rietschel E, Exner M, Lange L. 2009. Childhood hypersensitivity pneumonitis associated with fungal contamination of indoor hydroponics. *Int J Hyg Environ Health*. 212(1):18–20. doi: [10.1016/j.ijheh.2008.01.001](https://doi.org/10.1016/j.ijheh.2008.01.001).
- Engstad CS, Engstad RE, Olsen J-O, Osterud B. 2002. The effect of soluble  $\beta$ -1, 3-glucan and lipopolysaccharide on cytokine production and coagulation activation in whole blood. *Int Immunopharmacol*. 2(11):1585–1597. doi: [10.1016/s1567-5769\(02\)00134-0](https://doi.org/10.1016/s1567-5769(02)00134-0).
- Frankel M, Bekö G, Timm M, Gustavsen S, Hansen EW, Madsen AM. 2012. Seasonal variation of indoor microbial exposures and their relations to temperature, relative humidity and air exchange rates. *Appl Environ Microbiol*. 78(23):8289–8297. doi: [10.1128/AEM.02069-12](https://doi.org/10.1128/AEM.02069-12).
- Frankel M, Hansen EW, Madsen AM. 2014. Effect of relative humidity on the aerosolization and total inflammatory potential of fungal particles from dust-inoculated gypsum boards. *Indoor Air*. 24(1):16–28. doi: [10.1111/ina.12055](https://doi.org/10.1111/ina.12055).
- Gorni RL, Harkawy AS, Lawniczek-Walczyk A, Karbowska-Berent J, Wlazlo A, Niesler A, Golofit-Szymczak M, Cyprowski M. 2016. Exposure to culturable and total microbiota in cultural heritage conservation laboratories. *Int J Occup Med Environ Health*. 29(2):255.
- Grabek-Lejko D, Tekiela A, Kasprzyk I. 2017. Risk of biodegradation of cultural heritage objects, stored in the historical and modern repositories in the Regional Museum in Rzeszow (Poland). A case study. *Int Biodeterior Biodegrad*. 123:46–55. doi: [10.1016/j.ibiod.2017.05.028](https://doi.org/10.1016/j.ibiod.2017.05.028).
- Guiamet P, Borrego S, Lavin P, Perdomo I, de Saravia SG. 2011. Biofouling and biodeterioration in materials stored at the Historical Archive of the Museum of La Plata, Argentine and at the National Archive of the Republic of Cuba. *Colloids Surf B Biointerfaces*. 85(2):229–234. doi: [10.1016/j.colsurfb.2011.02.031](https://doi.org/10.1016/j.colsurfb.2011.02.031).
- Hagiuda R, Oda H, Kawakami Y, Hirose D. 2022. Species diversity based on revised systematics of xerophilic Aspergillus section Restricti isolated from storage rooms and houses in Japan. *Biocontrol Sci*. 27(2):65–80. doi: [10.4265/bio.27.65](https://doi.org/10.4265/bio.27.65).
- Ilie DC, Caciora T, Ilies A, Berdenov Z, Hossain MA, Grama V, Dahal RK, Zdrinca M, Hassan TH, Herman GV, et al. 2023. Microbial air quality in the built environment—Case study of Darvas-La Roche Heritage Museum House, Oradea, Romania. *Buildings*. 13(3):620. doi: [10.3390/buildings13030620](https://doi.org/10.3390/buildings13030620).
- Jones DB. 1978. Therapy of postsurgical fungal endophthalmitis. *Ophthalmology*. 85(4):357–373. doi: [10.1016/s0161-6420\(78\)35664-5](https://doi.org/10.1016/s0161-6420(78)35664-5).
- Kavkler K, Gunde-Cimerman N, Zalar P, Demšar A. 2015. Fungal contamination of textile objects preserved in Slovene museums and religious institutions. *Int Biodeterior Biodegrad*. 97:51–59. doi: [10.1016/j.ibiod.2014.09.020](https://doi.org/10.1016/j.ibiod.2014.09.020).
- Kenny LC, Aitken RJ, Baldwin PEJ, Beaumont GC, Maynard AD. 1999. The sampling efficiency of personal inhalable aerosol samplers in low air movement environments. *J Aerosol Sci*. 30(5):627–638. doi: [10.1016/S0021-8502\(98\)00752-6](https://doi.org/10.1016/S0021-8502(98)00752-6).
- Kofoed VC, Campion C, Rasmussen PU, Møller SA, Eskildsen M, Nielsen JL, Madsen AM. 2024. Exposure to resistant fungi across working environments and time. *Sci Total Environ*. 923:171189.
- Kolmodin-Hedman B, Blomquist G, Sikström E. 1986. Mould exposure in museum personnel. *Int Arch Occup Environ Health*. 57(4):321–323. doi: [10.1007/BF00406187](https://doi.org/10.1007/BF00406187).
- Koul B, Upadhyay H. 2018. Fungi-mediated biodeterioration of household materials, libraries, cultural heritage and its control. In: Gehlot P, Singh J, editors. *Fungi their role in sustainable development: current perspectives*; Chapter 32. Singapore: Springer. p. 597–615.
- Lai PS, Sheehan WJ, Gaffin JM, Petty CR, Coull BA, Gold DR, Phipatanakul W. 2015. School endotoxin exposure

- and asthma morbidity in inner-city children. *Chest*. 148(5):1251–1258. doi: [10.1378/chest.15-0098](https://doi.org/10.1378/chest.15-0098).
- Lenart-Boron A, Wieckowska M, Pollesch J. **2016**. Changes in the concentration of microbial aerosol in the premises of the Jagiellonian University Museum during the Night of Museums (in Polish). *Opuscula Musealia*. 24:143–149.
- Madsen AM. **2006**. Airborne endotoxin in different background environments and seasons. *Ann Agric Environ Med.* 13(1):81–86.
- Madsen AM, Frederiksen MW, Bjerregaard M, Tendal K. **2020**. Measures to reduce the exposure of waste collection workers to handborne and airborne microorganisms and inflammogenic dust. *Waste Manag.* 101:241–249. doi: [10.1016/j.wasman.2019.10.023](https://doi.org/10.1016/j.wasman.2019.10.023).
- Micheluz A, Manente S, Tigini V, Prigione V, Pinzari F, Ravagnan G, Varese GC. **2015**. The extreme environment of a library: Xerophilic fungi inhabiting indoor niches. *Int Biodeterior Biodegrad.* 99:1–7. doi: [10.1016/j.ibiod.2014.12.012](https://doi.org/10.1016/j.ibiod.2014.12.012).
- Micheluz A, Sulyok M, Manente S, Krska R, Varese GC, Ravagnan G. **2016**. Fungal secondary metabolite analysis applied to Cultural Heritage: the case of a contaminated library in Venice. *WMJ.* 9(3):397–408. doi: [10.3920/WMJ2015.1958](https://doi.org/10.3920/WMJ2015.1958).
- Mittal M, Siddiqui MR, Tran K, Reddy SP, Malik AB. **2014**. Reactive oxygen species in inflammation and tissue injury. *Antioxid Redox Signal.* 20(7):1126–1167. doi: [10.1089/ars.2012.5149](https://doi.org/10.1089/ars.2012.5149).
- Museernes-Samråd. **2021**. Anbefalinger for klima (temperatur og relativ fugtighed) i museernes udstillinger og magasiner (in Danish). Slots-og-Kulturstyrelsen, editor. Denmark: Slots-og-Kulturstyrelsen.
- Natarajan K, Meganathan V, Mitchell C, Boggaram V. **2019**. Organic dust induces inflammatory gene expression in lung epithelial cells via ROS-dependent STAT-3 activation. *Am J Physiol Lung Cell Mol Physiol.* 317(1):L127–L140. doi: [10.1152/ajplung.00448.2018](https://doi.org/10.1152/ajplung.00448.2018).
- Nigam N, Dhawan S, Nair M. **1994**. Deterioration of feather and leather objects of some Indian museums by keratinophilic and non-keratinophilic fungi. *Int Biodeterior Biodegrad.* 33(2):145–152. doi: [10.1016/0964-8305\(94\)90033-7](https://doi.org/10.1016/0964-8305(94)90033-7).
- Niu M, Shen F, Zhou F, Zhu T, Zheng Y, Yang Y, Sun Y, Li X, Wu Y, Fu P, et al. **2020**. Indoor air filtration could lead to increased airborne endotoxin levels. *Environ Int.* 142:105878. doi: [10.1016/j.envint.2020.105878](https://doi.org/10.1016/j.envint.2020.105878).
- Okpalanozie OE, Adebusoye SA, Troiano F, Cattò C, Ilori MO, Cappitelli F. **2018**. Assessment of indoor air environment of a Nigerian museum library and its biodeteriorated books using culture-dependent and-independent techniques. *Int Biodeterior Biodegrad.* 132:139–149. doi: [10.1016/j.ibiod.2018.03.003](https://doi.org/10.1016/j.ibiod.2018.03.003).
- Park J, Cox-Ganser J, Rao C, Kreiss K. **2006**. Fungal and endotoxin measurements in dust associated with respiratory symptoms in a water-damaged office building. *Indoor Air.* 16(3):192–203. doi: [10.1111/j.1600-0668.2005.00415.x](https://doi.org/10.1111/j.1600-0668.2005.00415.x).
- Pasquarella C, Sansebastiano GE, Saccani E, Ugolotti M, Mariotti F, Boccuni C, Signorelli C, Fornari Schianchi L, Alessandrini C, Albertini R. **2011**. Proposal for an integrated approach to microbial environmental monitoring in cultural heritage: experience at the Correggio exhibition in Parma. *Aerobiologia*. 27(3):203–211. doi: [10.1007/s10453-010-9189-4](https://doi.org/10.1007/s10453-010-9189-4).
- Perez C, Tiné L, Verriele M, Locoge N, Becquart F, Lors C. **2024**. Strategy for the monitoring of fungal colonization in buildings-challenges and perspectives. *Int Biodeterior Biodegrad.* 189:105764. doi: [10.1016/j.ibiod.2024.105764](https://doi.org/10.1016/j.ibiod.2024.105764).
- Pinheiro AC, Sequeira SO, Macedo MF. **2019**. Fungi in archives, libraries, and museums: a review on paper conservation and human health. *Crit Rev Microbiol.* 45(5–6): 686–700. doi: [10.1080/1040841X.2019.1690420](https://doi.org/10.1080/1040841X.2019.1690420).
- Rasmussen PU, Frederiksen MW, Carøe TK, Madsen AM. **2023**. Health symptoms, inflammation, and bioaerosol exposure in workers at biowaste pretreatment plants. *Waste Manag.* 167:173–182. doi: [10.1016/j.wasman.2023.05.042](https://doi.org/10.1016/j.wasman.2023.05.042).
- Rasmussen PU, Phan HUT, Frederiksen MW, Madsen AM. **2021**. A characterization of bioaerosols in biowaste pretreatment plants in relation to occupational health. *Waste Manag.* 131:237–248. doi: [10.1016/j.wasman.2021.06.009](https://doi.org/10.1016/j.wasman.2021.06.009).
- Rojas TI, Aira M. **2012**. Fungal biodiversity in indoor environments in Havana, Cuba. *Aerobiologia*. 28(3):367–374. doi: [10.1007/s10453-011-9241-z](https://doi.org/10.1007/s10453-011-9241-z).
- Saridaki A, Glytsos T, Raisi L, Katsivela E, Tsiamis G, Kalogerakis N, Lazaridis M. **2023**. Airborne particles, bacterial and fungal communities insights of two museum exhibition halls with diverse air quality characteristics. *Aerobiologia*. 39(1):69–86. doi: [10.1007/s10453-022-09775-2](https://doi.org/10.1007/s10453-022-09775-2).
- Siqueira JP, Sutton DA, Gené J, García D, Wiederhold N, Guarro J. **2018**. Species of Aspergillus section Aspergillus from clinical samples in the United States. *Med Mycol.* 56(5):541–550. doi: [10.1093/mmy/myx085](https://doi.org/10.1093/mmy/myx085).
- Slimen A, Barboux R, Mihajlovski A, Moullarat S, Leplat J, Bousta F, Di Martino P. **2020**. High diversity of fungi associated with altered wood materials in the hunting lodge of “La Muette, Saint-Germain-en-Laye, France. *Mycol Progress.* 19(2):139–146. doi: [10.1007/s11557-019-01548-5](https://doi.org/10.1007/s11557-019-01548-5).
- Smith WM, Fahle G, Nussenblatt RB, Sen HN. **2013**. A rare case of endogenous Aspergillus conicus endophthalmitis in an immunocompromised patient. *J Ophthalm Inflamm Infect.* 3(1):1–4.
- Sterflinger K. **2010**. Fungi: their role in deterioration of cultural heritage. *Fungal Biol Rev.* 24(1–2):47–55. doi: [10.1016/j.fbr.2010.03.003](https://doi.org/10.1016/j.fbr.2010.03.003).
- Timm M, Hansen EW, Moesby L, Christensen JD. **2006**. Utilization of the human cell line HL-60 for chemiluminescence based detection of microorganisms and related substances. *Eur J Pharm Sci.* 27(2–3):252–258. doi: [10.1016/j.ejps.2005.10.006](https://doi.org/10.1016/j.ejps.2005.10.006).
- Vesper S, McKinstry C, Haugland R, Wymer L, Bradham K, Ashley P, Cox D, Dewalt G, Friedman W. **2007**. Development of an environmental relative moldiness index for US homes. *J Occup Environ Med.* 49(8):829–833. doi: [10.1097/JOM.0b013e3181255e98](https://doi.org/10.1097/JOM.0b013e3181255e98).
- Vitte J, Michel M, Malinovschi A, Caminati M, Odebode A, Annesi-Maesano I, Caimmi D, Cassagne C, Demoly P, Heffler E, et al. **2022**. Fungal exposome, human health and unmet needs: a 2022 update with special focus on allergy. *Authorea*. 77:1–19. doi: [10.22541/au.164690732.28621722/v1](https://doi.org/10.22541/au.164690732.28621722/v1).

- Wiszniewska M, Walusiak-Skorupa J, Pannenko I, Draniak M, Palczynski C. **2009**. Occupational exposure and sensitization to fungi among museum workers. *Occup Med (Lond)*. 59(4):237–242. doi: [10.1093/occmed/kqp043](https://doi.org/10.1093/occmed/kqp043).
- Zalar P, Graf Hriberšek D, Gostinčar C, Breskvar M, Džeroski S, Matul M, Novak Babič M, Čremožnik Zupančič J, Kujović A, Gunde-Cimerman N, et al. **2023**. Xerophilic fungi contaminating historically valuable easel paintings from Slovenia. *Front Microbiol*. 14:1258670. doi: [10.3389/fmicb.2023.1258670](https://doi.org/10.3389/fmicb.2023.1258670).
- Zielńska-Jankiewicz K, Kozajda A, Piotrowska M, Szadkowska-Stańczyk I. **2008**. Microbiological contamination with moulds in work environment in libraries and archive storage facilities. *Ann Agric Environ Med*. 15(1): 71–78.