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Original paper

Evaluating the role of the working environment on to skin and upper respiratory tract microbiota of museum workers

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Abstract

Aims: The aim of the study was the investigation of the composition of cultivable microbiota colonizing the skin and the upper respiratory tract mucosa of museum workers, in order to evaluate the potential occupational risk to their health and wellbeing.

Methods: This study was voluntarily attended by 50 workers of a Romanian museum, for which their hands, noses and throats were sampled using sterile swabs. The API tests were used to identify the β -hemolytic strains, and penicillin and the cefoxitin susceptibility test was performed for *Staphylococcus* sp. strains.

Results: This study showed that a high percentage of the museum workers presented changes in the normal oropharyngeal and skin cultivable microbiota, indicated by the presence of dysbiosis, β -hemolytic microorganisms, methicillin and penicillin resistant staphylococci. The identification step revealed species of staphylococci which showed high levels of resistance to penicillin and methicillin.

Conclusions: The highlighted opportunistic and antibiotic resistant bacterial strains may be a risk factor for workers. The work environment should be further investigated as a possible source of contamination, since some of the species identified in the present study were also identified in a former study, made in the same museum, showing the possibility that contamination has occurred at the workplace.

Keywords

Occupational diseases, work environment, health, workers, microorganisms, contamination, museums, artifacts.

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Introduction

Human microbial exposure is common in many work settings and could have serious consequences to the workers' health (LIEBERS V, RAULF-HEIMSOTH M, BRÜNING T, 2008). Museums are among the crucial institutions involved in preserving the cultural heritage and offer diverse working environments for different specialists (archivists, curators, conservators, storeroom workers and office staff), many of these media having shown a high microbial air contamination (SKÓRA J, GUTAROWSKA B, PIELECH-PRZYBYLSKA K, et al, 2015).

Inside museums, microbial contamination does not only threaten with biodegradation of cultural heritage artifacts, but also represents a health risk for workers (GÓRNY RL et al, 2016). Microclimate factors (humidity, temperature, air currents, etc.) and organic nutrients offered by museum objects (leather, paper, wood, etc.) offer appropriate conditions for microbial development. Moreover, new microorganisms are also introduced in the museum environment by visitors and museum workers (NIESLER A et al, 2010).

The statistics show that the number of employees in the cultural workplace is rising (EUROSTAT/ 2018), while health and safety at work are still a priority of the entire European Union. In Romania, in 2017, there were reported 762 museums and public collections – in which 6989 employees are working (INSR/2019). Eurostat data show that 7,9% of the workforce suffered from occupational health problems, of which 36% resulted in absence from work for at least 4 days (Eurostat, 2018a, 2018c). The negative effects can be huge, starting from absenteeism and high medical costs or insurance premiums or the presence of sick workers (with the risk of mistakes occurring during the execution) until early retirement and loss of qualified personnel. The European Agency for Safety and Health at Work (EU-OSHA) estimated 3,9% of the global total gross domestic product (GDP) and 3,3% of European GDP is spent on occupational diseases or injuries (EU-OSHA, 2019). This percentage varies from country to country.

In recent years, more and more studies on the microbial contamination in museums, heritage archives or libraries are being conducted. However, the scientific papers usually focus on microbial contamination of the museum or library collections themselves, while the workers' exposure is ignored (JANKIEWICZ KZ et al, 2008; GÓRNY RL et al, 2016).

Health threats in museums frequently occur due to fungal contamination of objects and air, causing different afflictions, from allergies to fungal infections or mycotoxicoses (VIEGAS S et al, 2018). Fungal infections can be lethal in immuno-deficient individuals (PFALLER MA; DIEKEMA DJ). Microfungi species, such as *Candida*, *Aspergillus*, *Penicillium*, *Alternaria*, *Cladosporium*, *Mucor*, *Rhizopus*, *Trichoderma*, *Fusarium*, *Acremonium*, *Scedoporum*, *Paecilomyces*, *Botrytis*, *Chrysonila* and *Aureobasidium*, (JANKIEWICZ, KZ, 2008; GUTAROWSKA B et al, 2012; PFALLER MA; DIEKEMA DJ), and

bacterial strains such as *Staphylococcus epidermidis*, *S. haemolyticus* and *Bacillus sp.* (HAIDUCU M et al, 2014) were often mentioned in studies that examined the presence of microorganisms in indoor air of museums, archives or libraries. Epidemiological studies claim that healthy individuals develop sensitivity if they are exposed for a long time to different microbial contaminants (KALWASINSKA A; BURKSANDRA A; WILK I, 2012).

Studies on the seasonal monitoring of bacteriological and fungal contamination of museums work environment are few, and however, they usually focus on microbial contamination of the museum collections themselves, while the exposure of workers is ignored, potentially leading to inappropriate management of occupational health and safety issues. This is the first study reporting on the characterization of culturable skin and upper respiratory tract microbiota of museum employees in Romania.

Materials and Methods

The employees enrolled in this study are working in one of the most visited museums in Romania, located in the south of the Southern Romanian Carpathians, in a region with very rich vegetation.

Out of the total number of 140 employees, 50 agreed to participate in this study as volunteers.

Of these participating, 38 were women with average age of around 46 years and 12 were men, with an average age of 39 years.

All volunteers have signed an informed consent, stating the purpose of the study in which they participated; the sampling procedures; their rights as volunteers; risks and/or possible discomforts; details of confidentiality of analytical results.

It is worth mentioning that only 36 of them accepted to give all three types of samples (nose, throat, hands). The swab samples were collected using standardized protocols used in the clinical laboratory. The swabs were transported to the laboratory in Amies transport medium, cultivated to blood agar and incubated in aerobic and anaerobiosis conditions for 24 hours at 37°C. All samples were analyzed for assessing the number of the developed colonies and the hemolysis reaction. After Gram staining and microscopic visualization, the catalase test was performed.

All beta-hemolytic and catalase-positive colonies revealing Gram-positive cocci were tested for penicillin 10IU and cefoxitin (30 µg) susceptibility following the CLSI recommendations. The catalase-negative colonies revealing Gram-positive cocci were tested for agglutination with BioMerieux Slidex Staph-Kit. Depending on the microscopic aspects, API identification tests were used for the rest of the bacterial strains.

Results

The purpose of this study was to investigate the culturable skin and respiratory tract microbiota of museum workers in order to evaluate the influence of the working environment on the health of museum workers.

The analyzed parameters for an abnormal microbiota were: **i)** presence of dysbiosis, defined by the recovery of one single colony type from the analyzed samples;

ii) the presence of β -haemolytic colonies; **iii)** the presence of fungal colonies. The results obtained are presented in Figure 1.

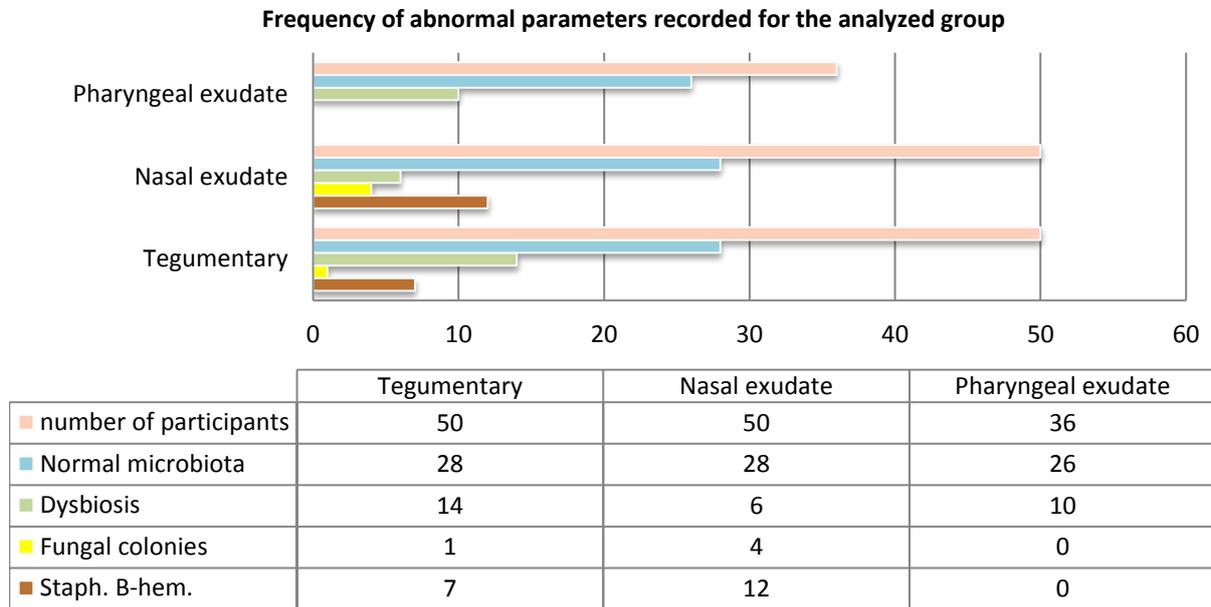


Figure 1. Study to investigate the culturable skin and respiratory tract microbiota of museum workers.

From the 50 volunteers, only 28 (56%) exhibited had normal results following the skin and nasal swabs analysis, while from the 36 participants agreeing to give throat swabs, 26 (72.2%) had a normal aspect of cultivable microbiota. From the investigated parameters, dysbiosis was the most frequent abnormal condition encountered in

the study, followed by the presence of β -haemolytic colonies.

Of the 27 isolated haemolytic *Staphylococcaceae* strains (Gram-positive cocci, catalase-positive), 24 of them showed resistance to penicillin and/or cefoxitin (Table 1), indicating the high rate of methicillin resistance in the analyzed population group.

Table 1. Antibiotic susceptibility testing of *Staphylococcaceae* strains

Antibiotics	CFU*
Resistance to Penicillin and Cefoxitin	20
Resistance to Cefoxitin, sensitive Penicillin	4
Sensitive to Penicillin and Cefoxitin	3

* CFU – Colony Forming Units

API tests did not reveal the presence of *Staphylococcus aureus*, but other species have been identified, such as: *Staphylococcus xylosum*, the most frequent species (62%), followed by *Staphylococcus haemolyticus* (21%), *Staphylococcus epidermidis* (13%) and *Staphylococcus sciuri* (4%). These strains have been mainly identified from nasal exudates, while *S. epidermidis* was identified only in skin samples.

The density of skin microorganisms is normally about 10^7 /cm² (DIMA MB et al, 2018). It has been shown that skin microbiota dysbiosis can be driven by the overgrowth of a certain species. For example, a US study performed in 2016 (DIMA MB et al, 2018) showed that commensal Gram-negative bacteria from healthy individuals are inhibited by the growth of Methicillin-Resistant *Staphylococcus aureus*.

In the present study, even though there was a large number of participants with dysbiosis, *Staphylococcus*

aureus was not isolated. However, other *Staphylococcus* sp. strains were isolated and many of them were resistant to antibiotics, as for example *Staphylococcus sciuri*. There are studies that claim the PBP2a protein from *S. sciuri* has 88% amino acid sequence identity to methicillin-resistant *S. aureus* PBP2a (ZEMAN M et al, 2017).

Only a small percentage of the tested workers harbored fungi on skin or nasal mucosa (1 versus 4).

The findings in this study can be correlated with findings from a former study that was conducted within a similar setup, in 2017, as part of the TEXLECONS project. Out of the four seasonal measurements, the maximum values recorded for mesophilic microorganisms were 1.4×10^4 CFU/m³, for hemolytic bacteria 350 CFU/m³ and for fungi up to 3×10^3 CFU/m³ (SCARLAT I et al, 2017). Some of the bacterial strains identified on the workers skin and mucosa have been also isolated from the museum environment, hence suggesting that the workplace may indeed

be contaminated by artifacts and indoor air, thus raising the need to further investigate the artifacts and indoor air within the workplace as a possible source of contamination.

Conclusions

This study demonstrates that microorganisms present in the work environment of museums can have effects on the health of workers. The results obtained show that more than half of the volunteer workers have values deviate from the reference values in at least one of the analyses, represented by the presence of dysbiosis, β -hemolytic microorganisms, methicillin and penicillin resistant staphylococci and, less frequently fungal colonization of skin and nasal mucosa. The work environment should be further investigated as a possible source of contamination, since some of the species identified in the present study were also identified in a former study, conducted in the workplaces of the investigated persons, showing the possibility that contamination has occurred at the workplace.

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