

## Evaluation of the Levels and Quality of Microbial Contamination in Medical Emergency Departments in Comparison to Other Workplaces

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### Abstract

Work in Hospital Emergency Departments (HEDs) exposes both the emergency ward staff and patients to infectious and in other way harmful biological agents. The results of this study shows the presence of pathogenic bacteria isolated by three different methods. It revealed 9.8% of pathogens detected by imprint method, 10.5% of pathogens by swabbing method, 17.6% and 22% in HEDs corridors and rooms, respectively, by air sampling method. In control workplaces (offices) pathogenic bacteria reached the level of 6.5% and 14.7% by imprint method and swabbing, respectively. The relatively low level of contamination by bacteria in HEDs may depend on the effectiveness of Standard Protective Precautions in the studied hospitals.

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**Key words:** harmful biological agents, Hospital Emergency Department, antimicrobial resistance, microbiological monitoring surface and air samples

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In accordance with the Act of 8 September 2006 on State Emergency Medical Services, the Hospital Emergency Department (HEDs) is a hospital ward whose main task is to provide comprehensive health services to adults and children in cases of sudden or life-threatening injuries or illnesses (isap.sejm.gov.pl) (ISAP, 2011). HEDs activities are focused mainly on stabilizing the patients, providing an initial diagnosis and transporting the patients to specialised wards for further treatment.

The character of HEDs work exposes both the emergency ward staff and patients to the presence of dangerous biological agents; however, the level of microbiological risk is not only related to the specifics of HEDs' functioning but it is also to the manner of performing the medical procedures by healthcare professionals. Followed correctly and in accordance with sanitation standards, medical procedures ensure security for both patients and medical personnel at the initial stage of treatment. The level of microbiological safety is further enhanced by ensuring that hospital environment is free of microorganisms capable of causing infection.

Pathogenic microorganisms in hospital environments such as HEDs are located generally on the surfaces of medical devices, as well as on the surfaces in direct contact with patients (walls, bed frames, medical devices switches, sinks *etc.*) and may pose the risk of hospital acquired infections (HAIs). The most common routes of transmission of potential pathogens are: direct contact with infected hospital personnel and contact between patients (cross transmission). Inaccurate cleaning of rooms or incorrect disinfection of medical equipment contributes to the spreading of pathogens through the devices routinely used in diagnosis, treatment and rehabilitation (Fiedotow and Denys, 2006). Bacteriological monitoring of hospital environment enables detection and differentiation of the colonizing and infective bacterial flora and provides the basis for effective empiric antibiotic therapy and eradication of the microorganisms from the ward environment (Maszkiewicz, 2007).

Systematic surveillance of infections, mainly by identifying the etiological agents of HAIs, enables identification of the most commonly found microorganism,

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posing a threat not only to the patient but also to the whole hospital environment (Heczko and Wójkowska-Mach, 2009).

This study used a random sampling strategy in HEDs environment and in a control workplaces (offices) not exposed to contact with harmful biological agents. The samples were taken from hospital surfaces (floors, walls, tables) and irregular, hard to access or frequently used surfaces (ventilation grilles, keyboards, medical device buttons, pens *etc.*) that may pose a risk of possible transfer of pathogenic microorganisms.

Environmental samples for this research were collected in the autumn and winter of 2014. All materials were sampled from areas with normal work system in HEDs in the presence of patients, visitors and medical personnel. Sampling apparatus was placed at height of around 0.8–1.2 m in a representative location of the examined spaces. The examined areas were varied in relation to the capacity and numbers of beds. All rooms were equipped in gravity ventilation systems. In total, 90 samples were collected in 10 Hospital Emergency Departments, including 20 air samples (HED's rooms and corridors) and 30 imprint samples; the last were taken in order to assess the total number of microorganisms. The remaining 40 samples were collected by swabbing high risk surfaces, *e.g.* door handles, keyboards, badges, pens. A similar sampling algorithm was adopted in the analysis of microbiological contamination of control workplaces, where a total of 80 samples were collected, 70 samples from surfaces (30 – imprints, 40 – swabs) and 10 air samples. In total, 170 environmental samples were collected from HEDs and offices. The biological material from imprinting samples Count-Tact® 3P™ Agar-CT3P (bioMérieux, France), together with samples collected using swabs, was inoculated on selective culture media and prepared in accordance with the generally accepted microbiological diagnostic procedures.

Air sampling was carried out with the assistance of a Coriolis Recon apparatus (Bertin, France); this technic collects a large amount of biological particles (0.5–10.0 µm) in a liquid sample at a flow rate of 6 m<sup>3</sup> (for 10 min) and allows preparation of a highly representative sample for further microbiological analysis.

Air samples were filtered through a 0.22 µm filter (Merck Millipore, Poland) using the filtering apparatus vacuum pump P504 Millipore (Merck Millipore, Poland). The filter was placed on the surface of Columbia agar medium with 5% sheep blood (Graso Biotech, Poland), and the culture of the biological material was incubated at 37°C for 24 h; the following stages were carried out in accordance with the generally accepted diagnostic scheme. Identification and testing the strains of bacteria for susceptibility to various groups of antibiotics was carried out using an automated system

Vitek 2 Compact (bioMérieux, France) following the manufacturer's instructions. For confirmation of antimicrobial resistance of bacterial isolates the disc diffusion method was used according to actual EUCAST recommendation ([www.korid.edu.pl/spec-rekomendacje-eucast.php](http://www.korid.edu.pl/spec-rekomendacje-eucast.php)). Controls for the study were conducted using following reference strains: *Escherichia coli* ATCC 25922, *Enterococcus faecalis* ATCC 29212, *Staphylococcus aureus* ATCC 29213, *Pseudomonas aeruginosa* ATCC 27853, *Klebsiella pneumoniae* ATCC 700603, *Streptococcus pneumoniae* ATCC 49619.

Statistical evaluation of the obtained results by three methods was done by Mann-Whitney test because the analysed data have not passed Shapiro-Wilk normality test (GraphPadPrism, USA). The difference is statistically significant at  $P < 0.05$ .

Harmful biological agents are a significant problem in occupational medicine and environmental health. Identification and characterization of the agents in the hospital environment makes it possible to conduct a reliable risk assessment of biological hazards to medical emergency staff in HEDs (Kramer *et al.*, 2006). The standards of assessment of the microbiological hygiene status of contact surfaces in medical institutions were suggested by the US Department of Agriculture ( $< 5$  cfu/cm<sup>2</sup> and  $< 1$  cfu/cm<sup>2</sup> for indicator organisms: *S. aureus*, *Clostridium difficile*, vancomycin-resistant *Enterococcus* (VRE), multidrug resistance (MDR) Gram-negative *Bacilli*, *Salmonella* spp., *E. coli* O157) (Dancer, 2004). According to these standards, the status of contact surfaces contamination (floors, walls, tables) in the examined HEDs falls within the permissible norm (Table I). All examined surfaces in HEDs *vs.* offices, except floors, revealed statistically significant differences. In similar studies concerning the levels of biological contamination of small scale medical equipment *e.g.* stethoscope (Shiferaw *et al.*, 2013), a significant degree of micro-contamination was revealed (micro-contamination  $\geq 20$  cfu per membrane; the equivalent of approx. 2 cfu/cm<sup>2</sup>). In the light of the above, the hygiene status of the examined contact surface in HEDs seems to be satisfactory. It may be the result of correct implementation of the Standard Protective Precautions and effective disinfection and decontamination procedures (Dancer, 2004). Microbiological contamination of air in HEDs rooms and corridors was on average at the level of  $2.1 \times 10^1$  cfu/m<sup>3</sup> and  $3.8 \times 10^1$  cfu/m<sup>3</sup>, respectively. The mean value of air micro-contamination in offices was  $1.4 \times 10^1$  cfu/m<sup>3</sup> (bacteria) (Table I).

According to the research of the past few decades, micro-contamination of air in hospital rooms varies depending on the sterility class of the examined hospital rooms *e.g.* from  $7.0 \times 10^1$  cfu/m<sup>3</sup> for bacteria and fungi as the highest acceptable concentration in

Table I  
Total number of bacteria from tested sites

| Tested sites                      | Imprint samples (cfu/25cm <sup>2</sup> )      |                       |           |   |                       |           | P value** |
|-----------------------------------|---|-----------------------|-----------|---|-----------------------|-----------|-----------|
|                                   | HEDs; n* = 30                                 |                       |           | Offices; n* = 30                              |                       |           |           |
|                                   | Mean values ± SD                              | Median values         | Min.–Max. | Mean values ± SD                              | Median values         | Min.–Max. |           |
| Tables                            | 8.7 × 10 <sup>1</sup> ± 8.4 × 10 <sup>1</sup> | 5.5 × 10 <sup>1</sup> | 18 – 260  | 2.6 × 10 <sup>1</sup> ± 2.9 × 10 <sup>1</sup> | 1.3 × 10 <sup>1</sup> | 2 – 90    | 0.0147    |
| Floors                            | 1.3 × 10 <sup>2</sup> ± 1.0 × 10 <sup>2</sup> | 1.1 × 10 <sup>2</sup> | 16 – 300  | 8.4 × 10 <sup>1</sup> ± 9.8 × 10 <sup>1</sup> | 4.4 × 10 <sup>1</sup> | 4 – 300   | 0.1502    |
| Walls                             | 3.5 × 10 <sup>1</sup> ± 3.4 × 10 <sup>1</sup> | 1.9 × 10 <sup>1</sup> | 6 – 100   | 1.0 × 10 <sup>1</sup> ± 7.5 × 10 <sup>0</sup> | 1.0 × 10 <sup>1</sup> | 0 – 20    | 0.0310    |
| <b>Total</b>                      | 8.6 × 10 <sup>1</sup> ± 8.6 × 10 <sup>1</sup> | 6.2 × 10 <sup>1</sup> | 6 – 300   | 4.0 × 10 <sup>1</sup> ± 6.6 × 10 <sup>1</sup> | 1.6 × 10 <sup>1</sup> | 0 – 300   | 0.0009    |
| Air samples (cfu/m <sup>3</sup> ) |   |                       |           |   |                       |           |           |
| Air                               | HEDs; n* = 20                                 |                       |           | Offices; n* = 10                              |                       |           | 0.0365    |
|                                   | 2.8 × 10 <sup>1</sup> ± 1.9 × 10 <sup>1</sup> | 2.8 × 10 <sup>1</sup> | 2 – 70    | 1.3 × 10 <sup>1</sup> ± 1.2 × 10 <sup>1</sup> | 8.0 × 10 <sup>0</sup> | 3 – 42    |           |

\* n = number of samples \*\* – calculated with Mann-Whitney test

a neurosurgical wards to  $7.0 \times 10^2$  cfu/m<sup>3</sup> as the highest acceptable concentration in treatment rooms (Górny, 2004). Other examples of microbiological assessment analyses are presented by Rubino (1995), according to whom, the level  $0-2.3 \times 10^1$  cfu/m<sup>3</sup> is pronounced as very good and the value  $>3.75 \times 10^2$  cfu/m<sup>3</sup> marks a highly unsatisfactory level of microbiological contamination (Charkowska, 2003).

According to new surveys the trends of microbiological air contamination are for an improvement of quantitative and qualitative status of bio-aerosols of the hospital environments (Ekhaise *et al.*, 2008; Hoseinzadeh *et al.*, 2013; Mirzaei *et al.*, 2014).

Microbiological air quality is frequently monitored in sterile hospital rooms (where the risk of infection is highest) however, there is little data about air micro-contamination of hospital corridors. Some research was presented by Park *et al.* (2013), who registered the air microbiological contamination in hospital corridors at the level  $7.2 \times 10^2$  cfu/m<sup>3</sup> for the bacteria. The values found here are much higher than the ones registered in “clean rooms” (*e.g.* operation rooms or haematology, pulmonology, obstetrics and gynaecology wards). The hospital wards with the highest level of sterility such as surgical and transplant units present the microbiological contamination in the range from  $0.01-1 \times 10^1$  cfu/m<sup>3</sup> (Park *et al.*, 2013).

However, in our research, both HEDs rooms and corridors represent the microbiological contamination at the level of  $2.7 \times 10^1$  cfu/m<sup>3</sup>, which places them within the limits of accepted standards (Gołofit-Szymczak *et al.*, 2013). Also in the research conducted by Augustowska and Dutkiewicz (2006) concentrations of bacteria in the air assumed values from  $1.0$  to  $9.6 \times 10^1$  cfu/m<sup>3</sup>.

Relatively low concentrations of microorganisms detected in the hospital environment (contact surfaces and air) may result from improved levels of hygiene in hospitals due to infrastructure development (air conditioning and highly efficient HEPA filters) and

implementation of modern and highly effective sanitary procedures. Another preventive measure reducing microbiological contamination in the hospital environment is the implementation of hospital infection control teams. This study, carried out in the winter, achieved quantitative results similar to the values observed by other researchers who conducted their tests in the same season (NSI, 2009; 2010; 2011).

In assessing the microbiological quality of samples, this research found mainly non-fermentative Gram-negative *Bacilli* (NFGNB) and *Stenotrophomonas maltophilia*, naturally resistant to many broad-spectrum antibiotics (*e.g.*  $\beta$ -lactams including all carbapenems) and coagulase-negative *Staphylococci* (CoNS) with constitutive macrolide-lincosamide-streptogramin B (cMLS<sub>B</sub>) and macrolide-streptogramin B (MS<sub>B</sub>) phenotypes (Table II).

These bacteria may constitute a considerable risk of HAIs due to their resistance to a large number of antibiotics and antimicrobial therefore cause treatment problems; the bacteria may be classified as potentially pathogenic agents. In the examined contact and swab samples collected from HEDs, NFGNB were prevalent. In air samples, besides the mentioned NFGNB *Acinetobacter* spp. and *S. maltophilia* (mainly isolated from HEDs corridors) large numbers of CoNS with constitutive MLS<sub>B</sub> (especially in HEDs rooms) were found. In the tested air samples from HED corridors mainly NFGNB were found; in HEDs rooms CoNS with constitutive MLS<sub>B</sub> were isolated. In hospital offices contact and swab samples contained mainly CoNS; in the air samples no pathogens were detected (Table II). Comparison of the proportional presence of pathogenic bacteria isolated by three different methods revealed 9.8% of pathogens detected by imprint method and 10.5% of pathogens by swabbing method; and in relation to air sampling 17.6% and 22% for HEDs corridors and rooms, respectively. In offices pathogenic bacteria reached the level of 6.5% and 14.7% for imprint method

Table II  
The percentage of potential pathogens in relation to the number of all isolates

| Places  | Samples                  | NFGNB<br>(non-fermentative<br>Gram-negative <i>Bacilli</i> ) |      | CoNS<br>(coagulase-negative<br><i>Staphylococci</i> ) |     |
|---------|--------------------------|--|------|---|-----|
|         |                          | Species  | %    | Species   | %   |
| HEDs    | Imprints (n = 30)        | * <i>B. cepacia</i>  | 2.0  | <sup>1</sup> <i>S. saprophyticus</i>                  | 3.9 |
|         |                          | <i>S. maltophilia</i>  | 2.0  |   |     |
|         |                          | <i>A. lwoffii</i>  | 2.0  |   |     |
|         | Swabs (n = 40)           | <i>A. lwoffii</i>  | 2.8  | Not detected  |     |
|         |                          | <i>A. haemolyticus</i>                                       | 2.8  |   |     |
|         |                          | <i>A. radioresistens</i>                                     | 2.8  |   |     |
|         | Air (n = 10)<br>Corridor | <i>S. maltophilia</i>  | 10.0 | Not detected  |     |
|         |                          | ** <i>A. baumannii cplx</i>                                  | 5.0  |   |     |
|         |                          | <i>A. lwoffii</i>  | 5.0  |   |     |
|         | Air (n = 10)<br>Rooms    | Not detected   |      | <sup>2</sup> <i>S. haemolyticus</i>                   | 2.6 |
|         |                          | <sup>2</sup> <i>S. hominis</i>                               | 2.6  |   |     |
|         |                          | <sup>3</sup> <i>S. lentus</i>                                | 2.6  |   |     |
| Offices | Imprints (n = 30)        | <i>A. lwoffii</i>  | 11.4 | <i>S. hominis</i>                                     | 2.9 |
|         |                          | <i>A. baumannii</i>  | 2.0  | <i>S. haemolyticus</i>                                | 2.9 |
|         |                          |  |      | <i>S. warnerii</i>                                    | 2.9 |
|         | Swabs (n = 30)           | Not detected   |      | <i>S. warnerii</i>                                    | 6.6 |
|         |                          |  |      | <i>S. hominis</i>                                     | 6.6 |
|         | Air (n = 10)             | Not detected   |      | Not detected  |     |

\* isolate with ESBL resistance phenotype (alert-pathogen), \*\*multidrug resistance alert-pathogen, n – number of samples, <sup>1</sup> 1 isolate with MS<sub>B</sub> and 1 isolate with cMLS<sub>B</sub> and methicilin-resistant coagulase-negative *Staphylococci* MRCNS, <sup>2</sup> isolate with cMLS<sub>B</sub>, <sup>3</sup> isolate with MRCNS and cMLS<sub>B</sub>

and swabbing, respectively. In the air samples pathogenic microorganisms were not detected.

Widespread usage of antibiotics, both in hospitals and ambulatory treatments, has led to the selection of pathogens with varied phenotypes of antimicrobial resistance *i.e.* alert-pathogens. Among the pathogens isolated from contact and air samples collected in HEDs, this study detected two multidrug-resistant *Acinetobacter baumannii* (MRAB) and producing extended spectrum of  $\beta$ -lactamases *Burkholderia cepacia* (ESBL+) (Table II). According to the Ordinance of the Polish Ministry of Health, December 23<sup>rd</sup>, 2011 (List of alert pathogens, Attachment 1) these belong to alert pathogens (isap.sejm.gov.pl).

Literature concerning microbiological quality assessment discusses mainly research conducted in wards requiring long-term hospitalising *e.g.* intensive care, internal, haematology, urology, ophthalmology and surgical wards (Kępa *et al.*, 2012; Nourmoradi *et al.*, 2012; Paluchowska *et al.*, 2012a; 2012b; Guzek *et al.*, 2013; Hoseinzadeh *et al.*, 2013; Seweryn *et al.*, 2014). In this study, the bacteriological quality analysis of isolates from hospital environment (HEDs) correlates with the profiles of potentially pathogenic species determined by other researchers in similar tests. For example, in other studies

the main bacterial isolates responsible for microbiological contamination of contact surfaces in hospitals (floors, medical tables, couches, washbasins) were methicillin-resistant *S. aureus* (MRSA), VRE, *Clostridium difficile*, multidrug-resistant *Acinetobacter* spp., *Pseudomonas* spp., *Enterococcus* spp., and additionally Enterobacteriaceae ESBL+ (*E. coli*, *Enterobacter* spp., *Salmonella* spp. and *Klebsiella* spp.) (Kramer *et al.*, 2006; Garcia-Cruz *et al.*, 2012; Weber *et al.*, 2013; Seweryn *et al.*, 2014).

According to Paluchowska *et al.* (2012b) the largest proportion of HAIs caused by alert-pathogens is registered in intensive care wards, burns units, internal, haematological and surgical wards. The pathogens isolated most often were multi-drug resistant, NFGNB (mainly *A. baumannii* and *P. aeruginosa*) which are recognised as the most problematic to control and eradicate (Paluchowska *et al.*, 2012b). CoNS and *Enterococcus* spp. are the key factors in HAIs and they are mostly isolated from surgical, haematological and oncology wards (Guzek *et al.*, 2013; Nourmoradi *et al.*, 2012; Hoseinzadeh *et al.*, 2013). Additionally MRCNS and NFGNB isolated from urology wards may constitute sources of infection (Kępa *et al.*, 2012).

The results quoted above show that the diversity of microorganisms detected in this study is comparable

with the standard trends. In quality testing of control samples alert-pathogens were not detected; however, potentially pathogenic strains (mainly CoNS) were found. The relatively low level of contamination by bacteria in HEDs may attest to the effectiveness of implementing the standard protective precautions in the examined hospitals. In quality assessment of HEDs the main bacterial strains found were NFGNB and CoNS; the presence of the bacteria may result from person to person transmission or introduction of pathogens from outside the hospital.

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