

Original Research

Effect of Carbon Filter Usage Period on the Secondary Emission of Bioaerosols

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Abstract

Increased concentrations of bacteria and fungi in the air at the waste sorting plants and landfills pose health risks for employees as well as inhabitants of neighbouring areas. The current study presents the results of identification and quantification of the microorganisms present in the airflow leaving the carbon filters installed in the Solid Waste Utilization Plant in Pomeranian Voivodeship (Poland) in order to remove the odorous compounds from the sorting halls. Volatile organic compounds VOCs were identified by the sorption method using the solid Tenax TA sorbent and analysed using a thermal desorption-gas chromatography-mass spectrometry system (TD-GC/MS). Before activating the system for neutralizing odours, VOC concentrations in the sorting hall remained at the level of 12.1-15.2 mg/m³ and after activation of the neutralization system total concentration of VOCs in the sorting hall decreased and after the period of six months was 4.3 mg/m³. The microbiological analyses included the following microorganisms: psychrophilic and mesophilic bacteria from the coli group (*Enterobacteriaceae* family) and *Escherichia coli*, mold fungi and yeasts-like, Mannitol-positive and negative *Staphylococci*. Six months after the activation of the neutralization system, a marginal increase in all the studied microbiological parameters was observed, with the highest increase in Mannitol-positive *Staphylococcus* strains.

Keywords: bioaerosols, VOC, bacteria, fungi, carbon filters, odour substances

Introduction

Communal waste generated in residential areas and partly in other places (workplace, preschools, schools,

shopping centres, railway stations, airports, etc.) constitutes mixtures of food and its packages as well as materials used in households. The structure of waste depends on the year's seasons (seasonality), where huge amounts of compost materials are generated in the summer-autumn period and ash in the winter-spring season [1]. It usually takes a couple of days before this

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waste is utilized, and during this time especially food undergoes biodegradation processes. The household waste in the first stage is transferred to the sorting plants. During the sorting process several compounds are released into the air of the sorting halls, which belong to the group of volatile organic compounds (VOC). Among them methane, aromatic hydrocarbons (i.e., toluene, styrene, xylene), sulphur compounds (i.e., hydrogen sulphide, thiols), terpenes (i.e., α -, β -pinene, d-limonene, camphene, 3-carene), amines, alcohols (butanol), aldehydes, ketones (e.g., acetic acid, butyric acid) and volatile fatty acids were identified [2-6]. Most of these substances belong to the group of odour active compounds (OAC) and some of them are toxic. The employees of the utilization plants are mostly exposed to these compounds (via respiration, skin absorption); however, the odours are also usually burdensome to nearby residents. Other sources of VOCs include the burning of fuels such as gas, wood and kerosene. VOCs can be released from road bitumens and caustic effluents from bitumen production [7, 8]. VOCs including volatile sulfur compounds (VCSs) can be released from ground tire rubber and ground tire rubber after reclamation [9]. The odorous compounds are usually accompanied by the presence of microorganisms (including pathogens) and their fragments, which form bioaerosols. Biological factors pose a very serious health hazard to humans as some of them have allergenic or toxic properties [10]. They include bacteria, fungi, viruses, protozoa and compounds excreted by microorganisms such as: endotoxins, exotoxins, glucans, metabolites of fungi, mycotoxins, and allergens [11-14]. Inhalation or ingestion is the main route of exposure to the spread of fungi and bacteria. Microbial volatile organic compounds (MVOCs) or decomposing

volatile microbial products found in bioaerosols may contribute to the development of various diseases [15, 16]. Depending on the composition of the bioaerosol, it may cause simple irritations, allergic reactions and dangerous infectious diseases [10, 17]. The highest threat is posed by the components of bioaerosols transmitted by means of air-dust or air-droplet routes, which penetrate the human organism through the skin and mucous membranes and are ingested with food [18-20]. Fungi and bacteria are well-known allergens (e.g., *Penicillium* and *Aspergillus*) that can grow in many work environments (composting, municipal waste removal, etc.) [15, 21].

The problem of odour removal constitutes a topic of research conducted by several science centres proposing various technical and technological solutions. Fig. 1 presents the available and most common deodorization methods based on different physical and chemical phenomena. All of these methods have their advantages and disadvantages described in more detail [22-24], and some are still not fully assessed.

The main objective of this study was the identification and quantification of microorganisms present in the airflow leaving the carbon filters that were installed in the Solid Waste Utilization Plant localized in the Pomeranian region of Poland in order to remove the odorous compounds from the sorting halls and reduce environmental odorous emissions. Measuring and identifying the microorganisms took place after the installation of the carbon filters at the outlet air. Simultaneously, concentrations of VOCs and microorganisms in sorting halls were determined. This study was performed from November 2016 to January 2018 at the solid waste utilization plant in Pomeranian Voivodeship.

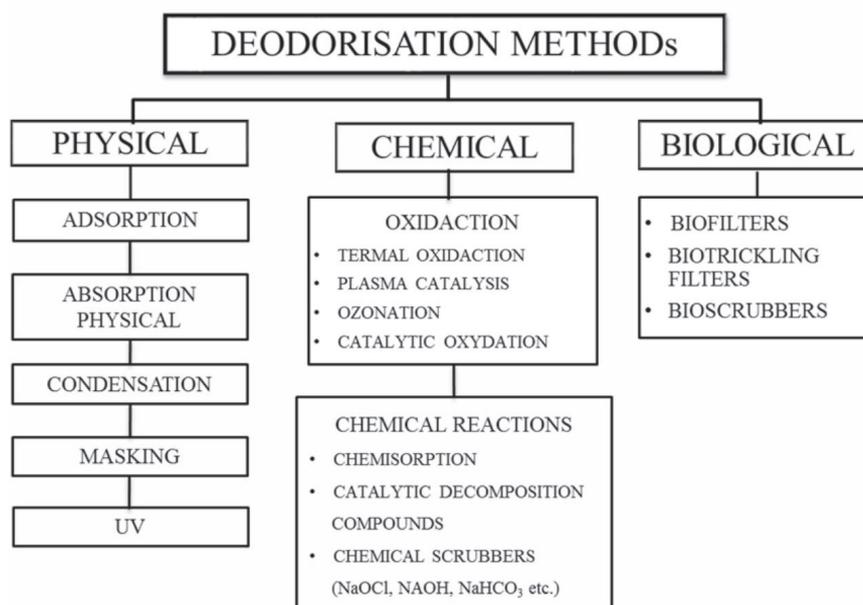


Fig. 1. Diagram of the most common deodorization methods.

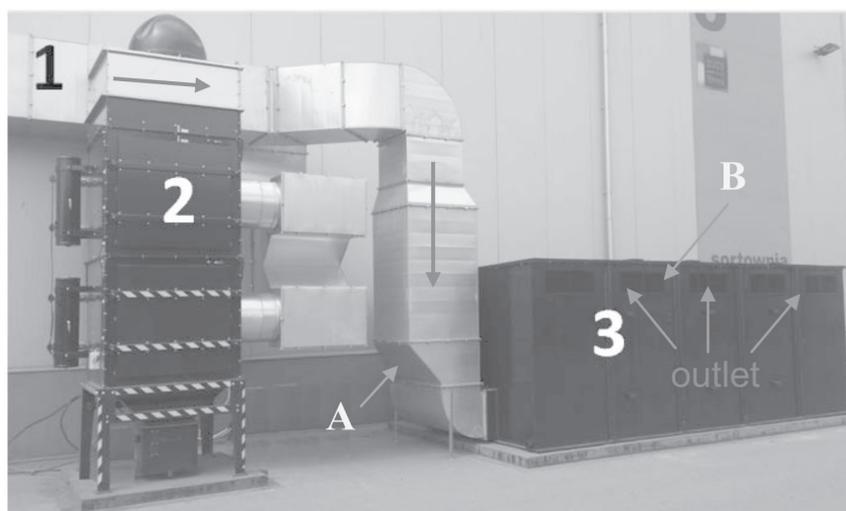


Fig. 2. Elements forming the neutralization system for the deodorization of air leaving the sorting hall with the direction of airflow: 1) ventilation canal used for the transportation of the airflow from the sorting hall, 2) cyclotron used for the removal of dust, and 3) neutralizer (container with the active carbon); locations of air sampling: A (air the coming carbon filters) and B (air leaving the carbon filters) (photo: A. Staniszewski).

Materials and Methods

Materials

As analytical standards the following chemicals were used: 1-bromo-4-fluorobenzene (purity 99.8%), 0.25 g, which was purchased in a solid form from dr. Ehrenstorfer GmbH (Germany). Methanol for gas chromatography (purity 99.9 %) was purchased from Merck KGAA (Darmstadt, Germany). A glass TD sample tube (1/4" × 90 mm) filled with 130 mg of Tenax TA (60-80 mesh) (Shimadzu, Japan) and capillary GC column Zebron (ZB-5ms), 30 m (length) : 0.25 mm (I.D.) : 0.25 μm (film), Phenomenex, (Shim-pol, USA) was used.

Instruments

Thermal desorption system TD-20 (Shimadzu, Japan) with sorbent Microtrap (Silcosteel tube 3.2-mm outer diameter (2-mm inner diameter) × 100 mm, filled with 60 mg Tenax TA) for refocusing of analytes prior to GC-MS analysis was used. GC-MS-QP 2010 ultra (Shimadzu, Japan).

Technical Description of the System Involved in the Neutralization of the Odorous Compounds

Locations of air sampling and the system used for the neutralization of the odorous compounds consists of three identical modules (N I, N II and N III) presented in Fig. 2. Arrangement of the ventilation in the sorting hall presented is Fig. 3. In one of the peripheral parts of the hall unsorted waste is delivered (the neutralization system III - this part is most-polluted and characterized by the largest odour emission), in the middle part (N II)

and the second peripheral part (N I) sorting takes place. Both parts are separated by a curtain that is not sealed, and the task is to physically divide the hall in order to improve the air condition and working conditions.

Determination of VOCs

VOCs were determined by the sorption method using the 130 mg of solid sorbent Tenax TA. 1.0 L and 0.6 L of air samples present in the sorting hall as well as 5.0 L of air samples subjected to neutralizer were collected. The samples were passed through glass tubes at the flow rate of 6.0 L/h. Then the tubes were closed with special caps and transported to the laboratory where on the same day they were analysed in the thermal desorption-gas chromatography-mass spectrometry system (TD-GC/MS). The quantitative determination of sum of VOC was carried using a single internal standard. 1-bromo-4-fluorobenzene (IS, 48.1 ng in 2 μL methanol) was added to each sorbent tube before the sampling process. All the concentrations are given in terms of added internal standard. According to literature data for Tenax TA, all quantified analytes have breakthrough volumes above the sampled air volume [25, 26]. Repeatability of standard deviation for such determination of the sum of VOC was less than 10% and is comparable with those published in literature.

Microbiological Studies

Air samples were also analysed using the Koch sedimentation method, where Petri dishes of a given diameter containing specific medium for the bacteria and fungi (selective or nutrient agar) were

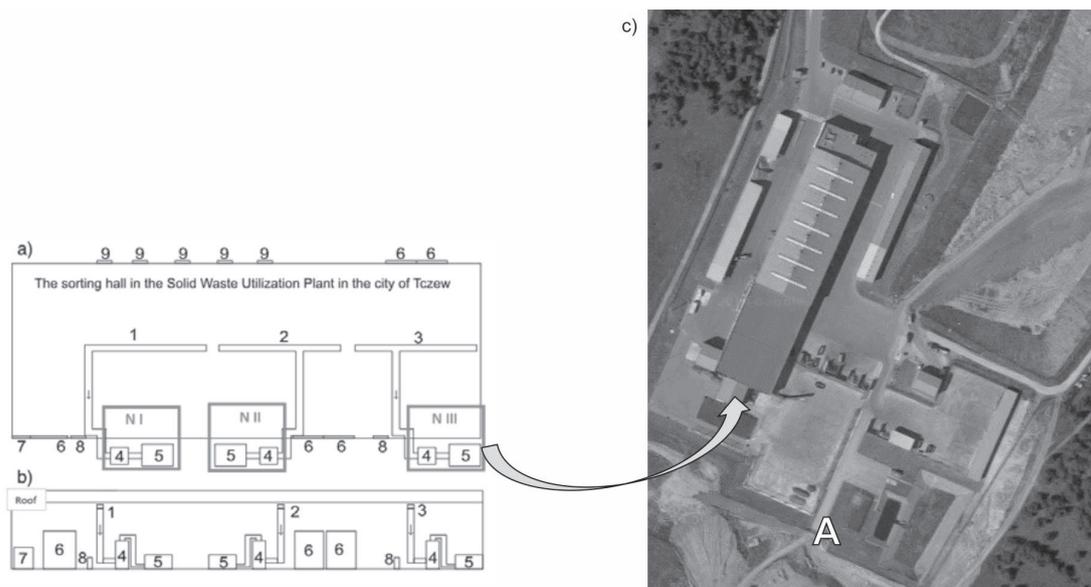


Fig. 3. Scheme of the ventilation in the sorting hall: a) view from the top, b) view from the side. 1. Ventilating canal of area number one, 2. Ventilating canal of area number two, 3. Ventilating canal of area number three, 4. Cyclotron used for the removal of dust, 5. Neutralizer (container with the active carbon), 6. Major entrance gate, 7. Minor entrance gate, 8. Entrance door, 9. Air-sucking device (fan), c) A- sampling point of the air masses flowing into the factory.

exposed for 5 to 10 minutes to the airflow in the sorting hall and leaving the carbon filters. After the incubation period of the samples, the total number of microorganisms in the air was determined using the Omelian formula expressed in the number of colony forming units (CFU) in 1 m³ of air. The microbiological analyses included the following microorganisms: psychrophilic and mesophilic bacteria from the coli group (*Enterobacteriaceae* family) and *Escherichia coli*, mold fungi and yeast-like, Mannitol-positive and negative *Staphylococci*. The total number of mesophilic bacteria was determined on trypticase soy agar (TSA) after incubation of the Petri dishes at 37°C for 24-48 hours. The total number of psychrophilic bacteria was also determined on the TSA, after the incubation of the Petri dishes at 22°C for 72 hours. The total number of *Staphylococcus* bacteria was determined in Chapman's medium after 24 hours incubation at 37°C. *Staphylococci* were identified as Manitol positive or negative strains. The ability to produce coagulase was also analysed in the test with the lyophilised rabbit plasma. The total number of the coli group (*Enterobacteriaceae* family) and *Escherichia coli* were determined on Chromocult Agar after incubation of the Petri dishes at 37°C for 24-48 hour. The total numbers of moulds and yeasts were determined on medium with chloramphenicol (YGC) after the incubation of the Petri dishes at 28°C for 7-10 days.

The identification of fungi was carried out using the Nikon Eclipse E2000 microscope on the basis of macro and microscopic features which were compared with the atlas of airborne fungal spores in Europe [27]. All the media used in the microbiological studies were purchased at Merck (Germany).

Microbiological studies were carried out three times – on 14 July 2017, 24 October 2017 and 25 January 2018. The average air temperatures on these days were 14°C, 6°C and 6°C, at relative humidity of 75%, 99% and 93%, respectively. Each time, control plates free of microbiological contaminants were also tested (unopened plates). In addition, on 25 January 2018, microbiological studies of the air masses flowing into the factory were performed (location of air sampling “A” in Fig. 3). The obtained results were similar to the results of the control, where traces of psychrophilic, mesophilic and Mannitol-negative *Staphylococci* and mold and yeast-like fungi were observed. Interpretation of the obtained results and microbiological assessment of the polluted air, according to the number of isolated microorganisms, was made on the basis of the Polish Standards (PN-Z-04111/02, 1989; PN-Z-04111/03, 1989). These standards currently do not apply, but so far have not been replaced by another document.

Statistical Analysis

Statistical analysis of obtained results was performed using Microsoft Excel 2010 (USA). Samples for testing microbiological parameters from the Solid Waste Utilization Plant were collected at different times after activation of the neutralization system. For VOC concentrations assessment samples were taken before starting the system, and during it working up to six months after system activation. Each sample collection period was performed in the same way. At each collection point after the carbon filters and at the sorting room samples were taken three times. To assess the effectiveness of the neutralization system

the calculated mean values were used. For each sample point, standard deviations were calculated to check the repeatability of measurements.

Results and Discussion

Characteristics of the VOCs

The system for the neutralization of odours was activated on 14 July 2017. Before activation of the neutralization system, VOC concentration levels in the sorting hall were in the range of 12.1-15.2 mg/m³ (mean: 13.6 mg/m³) (Fig. 4). After its activation the total concentration of VOCs in the sorting hall decreased, and after a period of six months achieved level 4.3 mg/m³ (mean: 6.5 mg/m³; the summation of individually determined compounds yielded a total VOCs value). The total concentrations of VOCs were about 39, 25 and 6 times lower for neutralization systems I, II and III, respectively. The differences in the VOC concentrations before the system for odour neutralization activation and after that were statistically significant ($p < 0.05$). In the VOC group, among others, the following compounds were identified: ethanol ($t_r = 1.56$ min), 1-methoxy-2-propanol and acetone ($t_r = 1.68$ min), acetic acid methyl ester ($t_r = 2.41$ min) and ethylbenzene ($t_r = 7.50$ min). Acetone was observed in the air in the coming carbon filters (air sampling a), while 1-methoxy-2-propanol was in the air leaving the carbon filters (air sampling b; Fig. 2). Acetone occurs on the concentrations of 0.004 mg/m³ (N I); 0.022 mg/m³ (N II); and 0.017 mg/m³ (N III). During the study, hydrogen sulphide and ammonia were also determined in the air of the sorting hall and at the outlet of the neutralizers using single-gas monitor GASBADGE pro (the European Standard EN50270 electromagnetic compatibility electrical apparatus for the detection and measurement of combustible gases, toxic gases or oxygen).

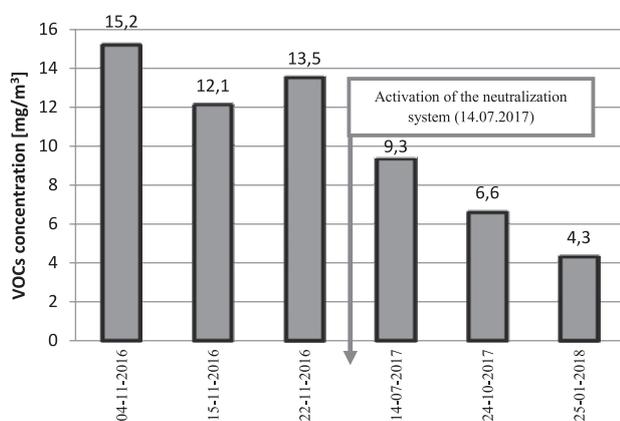


Fig. 4. Changes in the concentrations of VOCs in the applied chromatographic conditions.

Characteristics of the Cultured Bacteria

The results of this study show that the total number of the psychrophilic bacteria in the air in the sorting hall ranged from 3125 to 11250 CFU/m³. The total number of psychrophilic bacteria in the bioaerosol after the passage of air through neutralizer I ranged from 1574 to 2740 CFU/m³, after the passage through neutralizer II from 944 to 2646 CFU/m³, and after passage through neutralizer III ranged from 918 to 1858 CFU/m³. The total number of the mesophilic bacteria in the air in the sorting hall ranged from 1938 to 5180 CFU/m³. After the passage of the air through neutralizer I, the total number of mesophilic bacteria ranged from 1338 to 3540 CFU/m³, after the passage through neutralizer II ranged from 331 to 2457 CFU/m³, and after the passage through the neutralizer III ranged from 315 to 1638 CFU/m³. The number the *Enterobacteriaceae* family of bacteria in the air in the sorting hall ranged from 14 to 718 CFU/m³. The number of this bacteria in the bioaerosol after the passage of air through neutralizer I ranged from 0 to 158 CFU/m³, after passage through neutralizer II from 0 to 95 CFU/m³ and after passage through neutralizer III ranged from 0 to 7 CFU/m³. Including the number of *Escherichia coli* bacteria in the bioaerosol in the sorting hall ranged from 0 to 34 CFU/m³. After the passage through neutralizers I, II and III, the number of *Escherichia coli* bacteria ranged from 0 to 24 CFU/m³. The quantity of Mannitol-positive bacteria in the air samples from the sorting hall ranged from 94 to 307 CFU/m³ and the number of Mannitol-negative bacteria ranged from 188 to 384 CFU/m³. The number of this Mannitol-positive bacteria in the bioaerosol passing through the neutralizers I, II and III ranged from 158 to 881 CFU/m³, 79 to 763 CFU/m³ and 55 to 730 CFU/m³, respectively. The number of this Mannitol-negative bacteria in the bioaerosol after the passage of air through the N I ranged from 102 to 173 CFU/m³, after the passage through the N II from 70 to 173 CFU/m³ and after the passage through the N III ranged from 47 to 236 CFU/m³. Bacteria of the genus *Pseudomonas* spp. in the studied air were not isolated. Six months after the activation of the neutralization system, an increase in all the studied microbiological parameters was observed, with the highest increase in Mannitol-positive *Staphylococcus* strains (Fig. 5). In the air samples collected from the sorting hall, Mannitol-positive *Staphylococcus* (1.6%) and Mannitol-negative *Staphylococcus* (3.8%) – including *Staphylococcus epidermidis* (2.2%) and *Staphylococcus saprophyticus* (1.6%) – were isolated. From the air in the sorting hall the rods of the *Enterobacteriaceae* family (2.4%), including the *Escherichia coli* species (0.16%), were also isolated. The total number of psychrophilic bacteria isolated from the air in the sorting hall was 55.5% and the mesophilic bacteria 36.6%. In the air samples exiting

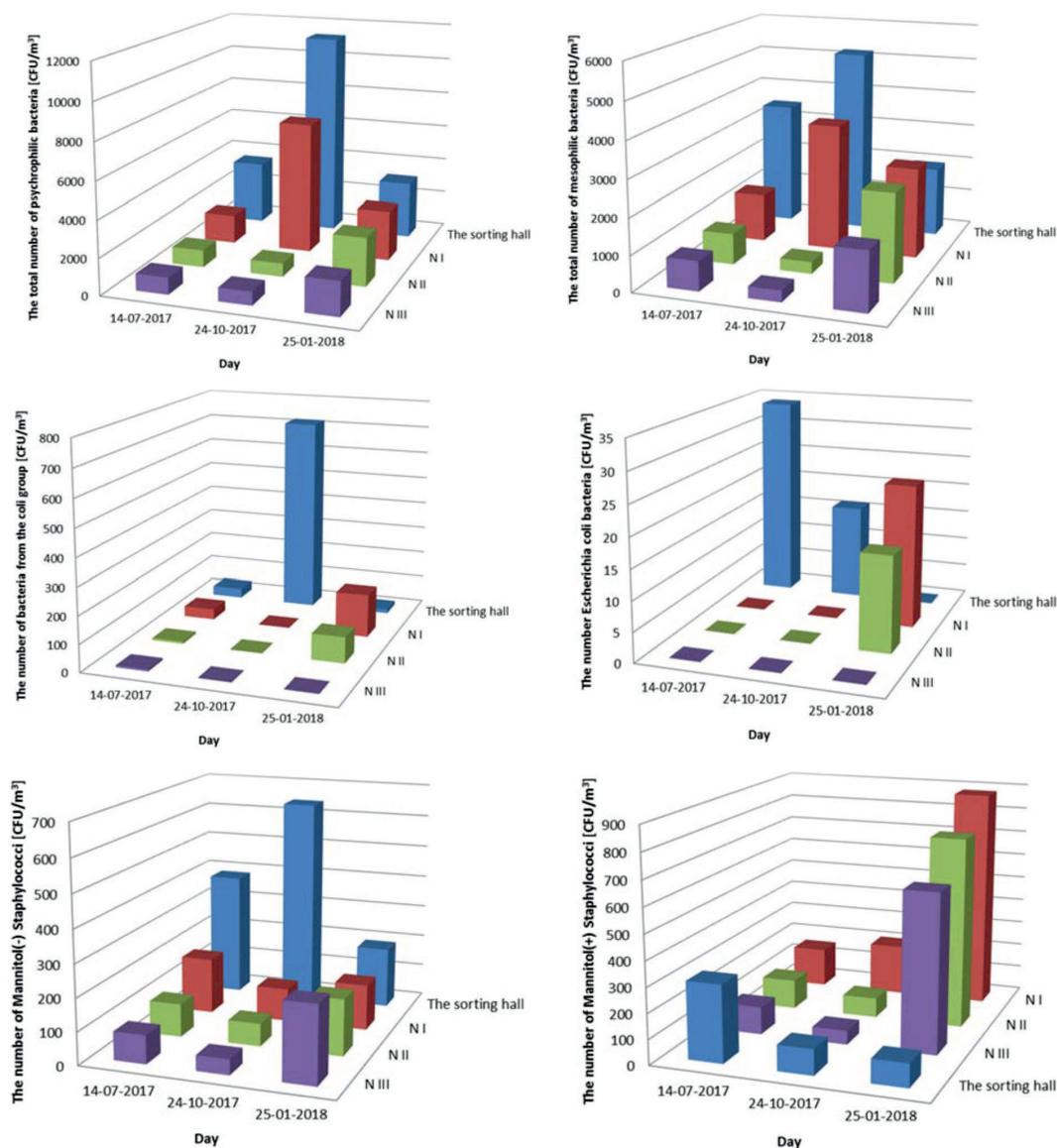


Fig. 5. Studied microbiological parameters (bacteria) during the research period.

carbon filters I, II and III (neutralizer), Mannitol-positive *Staphylococcus* (8.0%) and Mannitol-negative *Staphylococci* (3.0%), including *Staphylococcus epidermidis* (2.0%) and *Staphylococcus saprophyticus* (1.0%), were isolated. The rods of the *Enterobacteriaceae* family (0.8%), including *Escherichia coli* (0.1%) were also isolated. Bacteria of the *Pseudomonas* spp. genus in the studied air from the carbon filter neutralizer were not isolated. The total percentage of psychrophilic bacteria was 51.4% and the mesophilic bacteria 36.7%. The analysis showed significant differences in the concentrations of bacterial aerosol between the sorting room and the system of odour neutralization (NI-NIII; Table 1). The concentrations of bacterial aerosol were higher in the indoor air than in air after the carbon filters, the differences were statistically significant ($p < 0.05$). Six months after the activation of the neutralization system, a marginal increase in all studied microbiological parameters was observed

($p < 0.05$), with the highest increase in Mannitol-positive *Staphylococcus* strains (Table 2).

Characteristics of the Cultured Fungi

The quantity of mold and yeast-like fungi in the air samples from the sorting hall ranged from 273 to 2735 CFU/m³. The number of this fungi in the bioaerosol passing through neutralizers I, II and III ranged from 16 to 339 CFU/m³, 17 to 425 CFU/m³ and 8 to 245 CFU/m³, respectively (Fig. 6). From the collected air samples exiting carbon filters I, II, III (neutralizer) five species of mold fungi and one species of yeast *Rhodotorula rubra* (0.3%) were isolated. The most frequently isolated mold fungi was *Penicillium chrysogenum* (89.5%), *Aspergillus niger* (5.2%), *Aspergillus ochraceus* (2.4%) and *Mucor mucedo* (2.3%) were less frequently isolated. The least frequent in the studied air was the mold fungus *Alternaria alternata*

Table 1. Mean levels=standard deviation (SD) of microbial contamination in the sorting hall and after carbon filters (NI-NIII).

Type of microbial contamination	Psychrophilic bacteria [CFU/m ³] Mean±SD	Mesophilic bacteria [CFU/m ³] Mean±SD	Bacteria from the coli group [CFU/m ³] Mean±SD	<i>Escherichia coli</i> bacteria [CFU/m ³] Mean±SD	Mannitol(-) Staphylococci [CFU/m ³] Mean±SD	Mannitol(+) Staphylococci [CFU/m ³] Mean±SD	Mold and yeast-like fungi [CFU/m ³] Mean±SD
The sorting hall	5951.7±4591.9	3526.0±1622.0	255.7±400.5	16.7±17.0	404.3±227.2	167.7±120.7	2311.3±784.6
N I	3841.3±2975.0	2466.0±1101.9	65.7±82.3	8.0±13.9	138.7±35.6	414.7±404.5	186.7±162.3
N II	1441.0±1048.9	1218.0±1105.8	34.3±52.7	5.3±9.2	115.0±52.7	320.0±384.1	166.3±224.9
N III	1169.0±603.9	925.3±667.4	2.3±4.0	not detected	123.3±99.6	265.7±317.3	102.7±125.5

Table 2. Mean levels=standard deviation (SD) of microbial contamination in six months of activating the system of odour neutralization.

Type of microbial contamination	Psychrophilic bacteria [CFU/m ³] Mean±SD	Mesophilic bacteria [CFU/m ³] Mean±SD	Bacteria from the coli group [CFU/m ³] Mean±SD	<i>Escherichia coli</i> bacteria [CFU/m ³] Mean±SD	Mannitol(-) Staphylococci [CFU/m ³] Mean±SD	Mannitol(+) Staphylococci [CFU/m ³] Mean±SD	Mold and yeast-like fungi [CFU/m ³] Mean±SD
14-07-2017	1145.3±371.5	1009.0±285.7	18.0±18.2	not detected	120.7±45.9	128.7±25.7	105.7±86.0
24-10-2017	2891.3±3740.1	1395.3±1857.4	not detected	not detected	73.0±27.6	113.0±80.6	13.7±4.9
25-01-2018	2414.7±484.4	2205.0±492.0	84.3±79.5	13.3±12.2	183.3±48.3	758.0±125.6	336.3±90.0

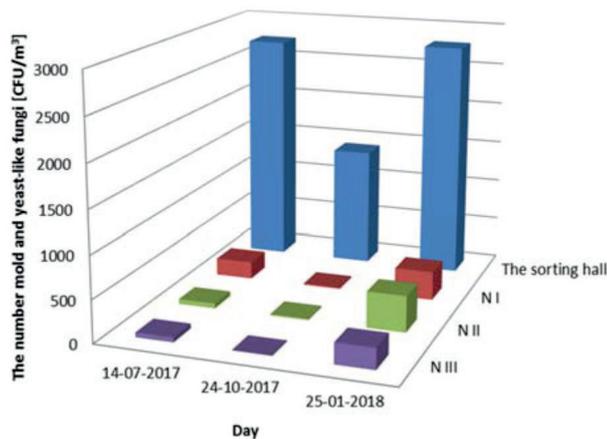


Fig. 6. Studied microbiological parameters (fungi) during the research period.

(0.3%). Table 3 shows the number of mold fungi and yeast-like fungi CFU/m³.

At present, there are no legal standards in Poland addressing the criteria and the normative values of the identification of microorganisms in polluted atmospheric air. There are also no world standards on the acceptable concentrations of biological contamination. In Poland the Interministerial Commission for Highest Concentration and Intensity of Agents Harmful to Health in the Working Environment proposes the acceptable concentrations of microorganisms in air at the level of 50000 CFU/m³ [28]. According to the World Health Organization (WHO), the acceptable value of microorganisms in air is 150 CFU/m³. Dutch experts believe that the fungal bioaerosol concentration over 10000 CFU/m³ becomes dangerous for human

Table 3. Numbers of mold fungi and yeast-like fungi CFU/m³

Study date	<i>Penicillium chrysogenum</i>	<i>Aspergillus niger</i>	<i>Aspergillus ochraceus</i>	<i>Mucor mucedo</i>	<i>Alternaria alternat</i>	<i>Rodotorhula rubra</i>
14.07.2017	2846	142	0	64	0	0
24.10.2017	1140	125	165	8	0	0
25.01.2018	3437	166	32	119	24	24
Sum [CFU/m ³]	7423	433	197	191	24	24
%	89.5	5.2	2.4	2.3	0.3	0.3

Table 4. Identification bacteria occurring on the premises and in the vicinity of municipal waste landfill sites in different countries.

Country	Location	Type of Bacteria	Ref.
Poland	Municipal landfills in Sosnowiec	<i>Micrococcus</i> spp., <i>Pseudomonas aeruginosa</i> , <i>Corynebacterium aquaticum</i> , <i>Staphylococcus xylosus</i> , <i>Staphylococcus sciuri</i> , <i>Staphylococcus auricularis</i> , <i>Pasteurella haemolytica</i> , <i>Bacillus cereus</i> , <i>Pseudomonas stutzeri</i> , <i>Rhodococcus</i> spp., <i>Sphingomonas paucimobilis</i> , <i>Bacillus amyloliquefaciens</i> , <i>Corynebacterium afermentans</i> , <i>Staphylococcus haemolyticus</i> ,	Lis et al. 2004 [11]
	Municipal Waste Disposal Complex at Zolwin-Wypaleniska near Bydgoszcz	<i>Pseudomonas fluorescens</i> , Actinomycetes, <i>Escherichia coli</i> , <i>Salmonella</i> spp., <i>Enterococcus</i> spp	Breza-Boruta 2012 [30]
Italy	An open composting facility flanked by a civil wastewater treatment plant located in a rural area of Northern Italy	<i>Mesophilic bacteria</i> , <i>psychrophilic bacteria</i> , <i>Pseudomonas</i> spp., <i>Enterobacteriaceae</i> , <i>Clostridium</i> spp.	Grisoli et al. 2009 [12]
Denmark	The waste in domestic areas in Copenhagen and containers to the trucks	<i>Acinetobacter</i> spp., <i>Bacillus</i> spp., <i>Cellulosimicrobium</i> spp., <i>Microbacterium</i> spp., <i>Micrococcus</i> spp., <i>Staphylococcus</i> spp., <i>Streptomyces</i> spp.	Madsen et al. 2016 [14]
Korea	Composting facility in Incheon City	<i>Bacillus amyloliquefaciens</i> , <i>Bacillus cereus</i> , <i>Staphylococcus lentus</i> , <i>Staphylococcus xylosus</i> , <i>Bacillus mycoides</i> , <i>Ochrobactrum anthropi</i> , <i>Streptomyces rochei</i>	Byeon et al. 2008 [18]
India	Municipal Solid Waste (MSW) aerobic compost treatment plant in Vidyaranyaapuram Mysore city)	<i>Klebsiella</i> spp., <i>Pseudomonas</i> spp., <i>Staphylococcus aureus</i> , <i>Enterobacter aerogenes</i> , <i>Salmonella</i> , <i>Bacillus</i> sp., <i>Escherichia coli</i> , <i>Flavobacterium</i> spp., <i>Staphylococcus xylosus</i>	Shyamala et al. 2014 [59]
Nigeria	Municipal solid waste dumpsite in Calabar metropolis	<i>Escherichia coli</i> , <i>Bacillus</i> spp., <i>Pseudomonas aeruginosa</i> , <i>Enterobacter</i> spp., <i>Klebsiella</i> spp., <i>Proteus</i> spp., <i>Salmonella</i> spp., <i>Staphylococcus aureus</i> , <i>Micrococcus luteus</i> , <i>Methanococcus</i> spp.	Bassey et al. 2016 [60]

Table 5. Identification fungi occurring on the premises and in the vicinities of municipal waste landfill sites in different countries.

Country	Location	Type of Fungi	Ref.
Poland	Municipal landfills in Sosnowiec	<i>Cladosporium herbarum</i> , Pink yeasts, White yeasts, <i>Arthrographis alba</i> , <i>Geotrichum candidum</i> , <i>Penicillium verrucosum</i> , <i>Staphylotrichum coccosporum</i> , <i>Phoma</i> sp., <i>Alternaria alternata</i> , <i>Cladosporium cladosporioides</i> , <i>Aspergillus niger</i> , <i>Myrothecium verrucaria</i> , <i>Humicola insolens</i> , <i>Trichoderma koningii</i> , <i>Aspergillus alutaceus</i> , <i>Aspergillus flavus</i> , <i>Aspergillus fumigatus</i> , <i>Fusarium dimerum</i> , <i>Fusarium solani</i> , <i>Gliocladium roseum</i> , <i>Malbranchea sulfurea</i> , <i>Rhizomucor pusillus</i> , <i>Thermoascus crustaceus</i> , <i>Acremonium charticola</i> , <i>Aureobasidium pullulans</i> , <i>Rhizopus rhizopodiformis</i>	Lis et al. 2004 [11]
	Municipal Waste Disposal Complex at Zolwin-Wypaleniska near Bydgoszcz	<i>Aspergillus Niger</i> , <i>Penicillium notatum</i> , <i>Rhizopus nigricans</i> , <i>Rhizopus oryzae</i> , <i>Sclerotinia sclerotium</i> , <i>Epicoccum nigrum</i> , <i>Arthrinium phaeospermum</i> , <i>Aspergillus fumigatus</i> , <i>Cladosporium herbarium</i> , <i>Alternaria alternata</i>	Breza-Boruta 2012 [30]
Italy	An open composting facility flanked by a civil wastewater treatment plant located in a rural area of Northern Italy	<i>Alternaria alternata</i> , <i>Arthrinium phaeospermum</i> , <i>Aspergillus chevalieri</i> , <i>Aspergillus clavato-nanica</i> , <i>Aspergillus flavus</i> , <i>Aspergillus fumigatus</i> , <i>Aspergillus glaucus</i> , <i>Aspergillus nidulans</i> , <i>Aspergillus niger</i> , <i>Aspergillus ochraeus</i> , <i>Aspergillus rescriptus</i> , <i>Aspergillus ruber</i> , <i>Aspergillus sydowii</i> , <i>Aspergillus terreus</i> , <i>Aspergillus thomii</i> , <i>Aureobasidium pullulans</i> , <i>Cladosporium cladosporioides</i> , <i>Cladosporium macrocarpum</i> , <i>Cladosporium oxysporum</i> , <i>Doratomyces</i> sp., <i>Epicoccum nigrum</i> , <i>Fusarium avenaceum</i> , <i>Fusarium lateritium</i> , <i>Fusarium poae</i> , <i>Fusarium sambucinum</i> , <i>Fusarium solani</i> , <i>Fusarium subglutinans</i> , <i>Mucor</i> spp., <i>Penicillium brevicompactum</i> , <i>Penicillium duclauxii</i> , <i>Penicillium minioluteum</i> , <i>Penicillium olsonii</i> , <i>Penicillium purpurogenum</i> , <i>Pseudorobillardaphragmitis</i> , <i>Rhizopus stolonifer</i> , <i>Scopulariopsis brevicaulis</i> , <i>Trichoderma viride</i> , <i>Trichothecium roseum</i>	Grisoli et al. 2009 [12]
Denmark	The waste in domestic areas in Copenhagen and containers to the trucks	<i>Aspergillus glaucus</i> , <i>Aspergillus nidulans</i> , <i>Aspergillus niger</i> , <i>Aspergillus versicolor</i> , <i>Cryptococcus magnus</i> , <i>Penicillium brevicompactum</i> , <i>Penicillium camemberti</i> , <i>Penicillium chrysogenum</i> , <i>Penicillium commune</i> , <i>Penicillium digitatum</i> , <i>Penicillium expansum</i> , <i>Penicillium glabrum</i> , <i>Penicillium italicum</i> , <i>Penicillium olsonii</i> , <i>Paecilomyces variotii</i> ,	Madsen et al. 2016 [14]
Korea	Composting facility in Incheon City	<i>Pseudallescheria boydii</i> , <i>Aspergillus fumigatus</i> , <i>Chaetomium</i> spp., <i>Penicillium</i> spp.,	Byeon et al. 2008 [18]
India	(Municipal Solid Waste (MSW) aerobic compost treatment plant in Vidyanarayapuram Mysore city)	<i>Alternaria</i> sp., <i>Aspergillus niger</i> , <i>Aspergillus fumigates</i> , <i>Aspergillus flavus</i> , <i>Aspergillus rapens</i> , <i>Fusarium</i> , <i>Mucor mucedo</i> , <i>Penicillium</i> sp., <i>Rhizopus stolonifer</i> , <i>Trichoderma</i> sp., <i>Varicosporium</i> ,	Shyamala et al. 2014 [59]
Nigeria	Municipal solid waste dumpsite in Calabar metropolis	<i>Candida albicans</i> , <i>Aspergillus</i> sp., <i>Saccharomyces</i> sp., <i>Candida tropicalis</i> , <i>Penicillium</i> sp., <i>Candida parapsilosis</i>	Bassey et al. 2016 [60]

health [29]. The authors of this study analysed the microbiological contamination of the atmospheric air based on: The Polish Standards (PN-89/Z-04111/02 and PN-89/Z-04111/03) and Report 12 of the CEC (Commission of the European Communities 1993). Indoor air quality and its impact on man. Biological particles in indoor environments. The average number of psychrophilic and mesophilic bacteria isolated from the air leaving the carbon filter was 2150 CFU/m³ and 1536 CFU/m³. The concentrations of these bacteria ranged from 1000 to 3000 bacteria in 1 m³, which according to the Polish Standard

PN-89Z-04111/02 is defined as moderately contaminated. According to the CEC, this was defined as highly and very highly contaminated. The average number of Mannitol-positive and Mannitol-negative *Staphylococci* bacteria in the sorting hall was 168 CFU/m³ and 333 CFU/m³, respectively, whereas the air leaving the carbon filters 630 CFU/m³ and 126 CFU/m³. The number of Mannitol-positive and Mannitol-negative *Staphylococci* exceeded acceptable values defined by the Polish Standard PN86/Z-04111/02. On 25 January 2018 the number of Mannitol-positive *Staphylococci* exceeded acceptable values by 35 times

Table 6. Review of literature on the concentrations of bacteria and fungi in the outdoor air.

Country	Location	Microorganisms concentration	Ref.
Poland	City Poznań	Total bacteria max: 13000 CFU/m ³ Mold fungi max: 9000-16000 CFU/m ³	Bugajny et al. 2005 [33]
	City Toruń	Bacteria heterotrophic mesophilic mean: 79-189 CFU/m ³ Family <i>Enterobacteriaceae</i> mean: 2-6 CFU/m ³ Mold fungi mean: 49-140 CFU/m ³ U m ³	Donderski et al. 2004 [43]
	City Bydgoszcz	Bacteria heterotrophic mean: 56-467 CFU/m ³ Staphylococci mean: 0-11 CFU/m ³ Mold fungi mean: 511-748 CFU/m ³	Małecka - Adamowicz et al. 2015 [44]
	City Kraków	Bacteria heterotrophic mesophilic: 11-327 CFU/m ³ Staphylococci mean: 10-69 CFU/m ³ , max: 863 CFU/m ³ Mold fungi mean: 226-111 CFU/m ³ , max 3447 CFU/m ³	Lenart-Boroń et al. 2014 [45]
	City Hel	Bacteria heterotrophic mesophilic max: 38 CFU/m ³ Mold fungi max: 600 CFU/m ³	Marks et al. 2001 [46]
	City Gdynia	Total bacteria max: 943 CFU/m ³ Mold fungi max: 266 CFU/m ³	Michalska et al. 2010 [47]
	City Sopot	Total bacteria max: 786 CFU/m ³ Mold fungi max: 2030 CFU/m ³	Michalska et al. 2010 [47]
Austria	City Graz	Bacteria heterotrophic mesophilic: 0-2500 CFU/m ³ Xerophilic fungi: 30-2300 CFU/m ³	Haas et al. 2013 [38]
Germany	City Kassel residents near composting site	Total bacteria: 22000-510000 CFU/m ³ Mold fungi: 7700-130000 CFU/m ³	Herr et al. 2003 [48]
Spain	Murcia	Total bacteria mean: 144 CFU/m ³ Mold fungi mean: 388 CFU/m ³	Soto et al. 2009 [49]
Greece	Thessalonki	Total bacteria mean: 120000 CFU/m ³	Genitsaris et al. 2017 [50]
Jordan	Al- Mafraq	Total bacteria max: 2055 CFU/m ³ Mold fungi max: 295 CFU/m ³	Jacob et al. 2016 [51]
China	Coastal region Qingdao	Total microorganisms: 85000-166000 CFU/m ³	Li et al. 2011 [52]
	City Xian	Total microorganisms: 77000-1421000 CFU/m ³	Xie et al. 2018 [53]
	City Beijing	Total bacteria: 71-22100 CFU/m ³ , mean: 2217 CFU/m ³	Fang et al. 2007 [54]
USA	City Cincinnati	Mold fungi: 0-3882 CFU/m ³	Lee et al. 2006 [34]
	New York City	Total microorganisms onshore winds mean: 580 CFU/m ³ Total microorganisms offshore winds mean: 778 CFU/m ³	Montero et al. 2016 [55]
Turkey	City Edirne	Total bacteria: 222-656 CFU/m ³ Staphylococci: 48-121 CFU/m ³	Aydogdu et al. 2010 [56]
India	City Gwalior	Total bacteria: 9000 CFU/m ³ Mold fungi: 2600 CFU/m ³	Yadav et al. 2015 [57]
	Maharashtra composting facility near Mumbai	Total bacteria mean: 3800-54000 CFU/m ³	Pahari et al. 2016 [58]

and the number of Mannitol-negative *Staphylococci* by 5 times. High concentrations of *Staphylococci* bacteria can influence air quality and pose a potential environmental risk [30]. The quantitative analysis of the air leaving the carbon filters showed the presence of *Enterobacteriaceae* in the range of 0 to 158 CFU/m³. The number of *Escherichia coli* bacteria ranged 0 to 24 CFU/m³. Epidemiological studies on the *Enterobacteriaceae* family in the air show that the highest concentrations of these bacteria were in wastewater treatments and on municipal waste landfills

[31, 32]. Identified bacteria occurring on the premises and in the vicinity of the municipal waste landfill site in different countries are comparable (Table 4). *Pseudomonas* spp., *Bacillus* spp., *Staphylococcus* spp., the family *Enterobacteriaceae* were most frequently identified. The positive conclusion of these results is the lack of high contamination of yeast-like fungi of the air leaving the carbon filter. The average number of mold and yeast-like fungi in the air from the sorting hall were 2311 CFU/m³ and in the bioaerosol passing through the carbon filters was 152 CFU/m³. According to the

Polish Standard PN-89Z-04111/03, air containing over 10000 CFU/m³ fungi poses a potential environmental risk. The concentrations of mold and yeast-like fungi in the air passing through the carbon filter points were relatively clean, and obtained values did not exceed 3000 CFU/m³, which according to the CEC is defined as moderately contaminated [33-35].

Identified fungi occurring on the premises and in the vicinity of the municipal waste landfill site in different countries are comparable (Table 5). The genus *Aspergillus* spp, *Cladosporium* spp, and *Penicillium* spp were most frequently identified.

A comparison of the number of microorganisms isolated from atmospheric air in different countries is presented in Table 6. The above-mentioned studies show that there is a large range in the number of bacteria in the atmospheric air. When comparing the concentrations of microorganisms in different countries, it should be remembered that such a diverse number of microorganisms in the air is affected by the source of emissions, distance from it, its size, survival of microorganisms, meteorological conditions, year season and the amount of dust suspended in the air [36-39]. The *Enterobacteriaceae* family includes, along with many harmless symbionts, many of the more familiar pathogens, such as *Salmonella* and *Escherichia coli*. Most members of this family constitute the gut flora found in the intestines among humans and other animals. Some enterobacteria produce endotoxins and active biological lipopolysaccharides (LPS), which are found in the outer membrane of Gram-negative bacteria and are released after the destruction of the cell wall [14, 16]. The most important effects of exposure to endotoxins are toxic pneumonitis (inhalation fever, organic dust toxic syndrome (ODTS), septic shock) [31,40].

Data from the literature point to higher concentrations of bacteria and fungi in the atmospheric air at the sewage treatment plant and municipal waste utilization plants (Table 4) than in this study. When comparing the amount of bacterial and fungal bioaerosol contamination in the study with the recommended standards of air leaving the carbon filters (neutralizer), it should be remembered that even if the concentration was lower than the reference values under certain circumstances, potentially pathogenic microorganisms may cause health problems [29, 40]. There are still no epidemiological data defining the relationship between exposure to a risk factor and the health effect caused by its action [20]. Adsorption and colonization of microorganisms on activated carbon is poorly understood.

The study conducted by Le Pape et al. [41] shows that the use of silver in activated carbon fibers can have a bactericidal effect against *Escherichia coli* and *Staphylococcus aureus*, and also fungicidal properties. The use of silver together with activated carbon can be a promising air-cleaning method [42].

Conclusion

This work shows the results of studies on the microbiological contamination and VOC concentrations in the air of the Solid Waste Utilization Plant. The quality of inhaled air has a significant impact on health and well-being of employees as well as for the inhabitants of neighbouring areas. Data from several studies point to microorganisms, and their secondary metabolites may pose toxic effects on the human respiratory system. Therefore, increased amounts of bacteria and fungi in the air should be the basis for monitoring the level of microorganisms around recycling plants. The degree of exposure of organisms to compounds from the VOC group depends to a large extent on the length of exposure and levels of analyte content that is inhaled. Data from several studies also point to prolonged exposure to these compounds possibly leading to permanent damage to the liver, kidneys and nervous system. Therefore, the reduction of emissions of these compounds to the environment is mandatory.

There is still little information in the world's literature about the possibility of colonization by bacteria and mold and yeast-like fungi on active carbon, which in many installations is part of the odor neutralization system. These microorganisms may be a source of secondary bioaerosol emissions to the atmosphere. The obtained results of identification studies and quantitative analyses of the microorganisms presented in the air stream leaving the systems of the carbon filters confirmed that the neutralization system effectively eliminates the odours from the sorting room. The active carbon layers reduced VOC concentrations in the air stream by up to 4.3 mg/m³, and at the same time decreased the concentration of odours in the sorting room. Six months after activating the neutralization system, an increase in all the studied microbiological parameters was observed. The highest increase in the number of Mannitol-positive *Staphylococcus* strains was detected, and the most frequently isolated mold fungi was *Penicillium chrysogenum* (89.5%).

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Conflict of Interest

The authors declare no conflict of interest.

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