



Assessment of Airborne Microorganisms in a Swine Wastewater Treatment Plant

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Abstract

Quantification of the airborne microorganisms (bacteria and fungi) at a swine wastewater treatment plant was performed. Microbial samples were collected at three different phases of the treatment process over a 1-yr period. Cultivation methods based on the viable counts of mesophilic heterotrophic bacteria and fungi were performed. The concentrations of airborne bacteria ranged up to about 5×10^3 colony-forming unit (CFU)/m³, and those of airborne fungi ranged up to about 9×10^2 CFU/m³. The primary treatment (e.g., screen, grit removal, and primary sedimentation) was found to be the major source of airborne microorganisms at the site studied, and higher levels of airborne bacteria and fungi were observed in summer. High levels of the respirable bioaerosol (0.65 to 4.7 μ m in size) were detected in the aeration phase. Among the environmental factors studied, temperature was strongly associated with fungal aerosol generation (with a Spearman correlation coefficient of 0.90 and p-value <0.01). Occupational biorisks are discussed based on the observed field data.

Keywords: Bioaerosol, Seasonal variation, Size distribution, Swine wastewater

1. Introduction

The rapid industrialization of livestock production has led to one of the world's most pressing environmental problems, including problems caused by the large quantities of wastewater generated [1]. Collection and treatment plants for wastewater have been used extensively to meet demands for lower effluent concentrations.

Wastewaters contain large amounts of pathogenic and non-pathogenic microorganisms, some of which may be a potential health hazards for workers [2]. Airborne release from these sites may be a critical pathway for pathogen movement off the sites [3]. Many studies have been carried out to evaluate the biological risks of aerosols emitted from municipal wastewater treatment plants [4-10]. Teixeira et al. [4] reported that in a municipal wastewater treatment plant, the indoor culturable bioaerosol concentration varied up to 5×10^4 colony-forming unit (CFU)/m³ for total bacteria, and 1.4×10^4 CFU/m³ for total fungi. The bioaerosol emission also varied according to the treatment stages [9, 11].

The exposure of workers to airborne microorganisms may vary depending on the type and capacity of the facility, the activities performed, and the weather conditions. Livestock-related facilities are a potential source of pathogenic bioaerosols [12],

especially *Campylobacter* spp. (responsible for campylobacteriosis), avian influenza virus, Newcastle disease virus, *Escherichia coli* (colibacillosis), *Salmonella* spp., and foot-and-mouth disease virus. Ko et al. [3] investigated bioaerosols released from twelve swine farms with different manure treatment technologies, using a liquid impinger for microbiological air sampling. They found that the concentration of airborne culturable bacteria ranged from 10^2 to 10^5 CFU/m³, and the airborne culturable fungal concentrations ranged from 10 to 10^3 CFU/m³. However, the liquid impinger had a 0.3 μ m cut-off diameter, which could result in lower collection efficiency for smaller-sized bioaerosols. The liquid impinger is less efficient for traditional cultivation-based detection compared to solid impactors, which are comparatively efficient for culturable bacterial sampling based on the results of molecular microbial analysis [13]. Since treated wastes differ at the various sites, the initial numbers and concentrations of microorganisms could be site-specific. Hence, further studies are required to characterize the bioaerosol emission in the swine wastewater treatment plants.

The environmental effects on bioaerosol emission in a swine wastewater treatment plant are not well known to the best of our knowledge. The size distribution of bioaerosol emitted from a plant is important to study, because the deposition of bioaerosol particles in the respiratory system depends on their aerodynam-

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Received July 23, 2012 Accepted November 28, 2012

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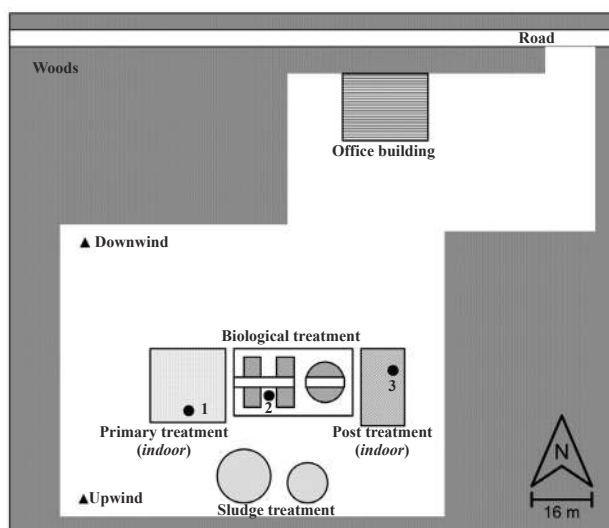


Fig. 1. Schematic diagram of the swine wastewater treatment plant and sampling sites indicated with (●): (1) primary treatment (indoor); (2) biological treatment; (3) post treatment (indoor). Upwind and downwind external control sites are indicated with (▲).

ic diameter, and the effects on human health are related to the sizes and physical properties of the particles [14].

This study has been conducted to examine the generation characteristics of airborne bacteria and fungi emitted from a swine wastewater treatment plant by using a solid impact air sampler. Cultivation methods based on the viable counts of mesophilic heterotrophic bacteria and fungi were performed. We characterized the seasonal change in airborne microorganism concentration at different stages of the treatment process (primary treatment, biological treatment, post-treatment). We also explored the distribution characteristics of airborne microorganisms for each particle size in each of the processes. The degree of correlation between airborne microorganisms and environmental conditions at the test sites was also determined.

2. Materials and Methods

2.1. Plant Description and Sampling Locations

The study was performed at the swine wastewater treatment plant, with activated sludge biological treatment, located in Hallym, Jeju Island, Korea. The selection of the study site was based on the structure of the plant (indoor and outdoor structures), and the plant was also located away from any existing sources of anthropogenic air pollution. A schematic diagram of the plant is presented in Fig. 1.

The plant receives 100 m³/day of wastewater, transported by truck and tank trailer, and generated by swine farms scattered all over Jeju Island. The plant has three separate phases of treatment process, consisting of primary treatment, biological treatment, and post treatment. Raw wastewater enters an indoor primary treatment process (hereafter referred to as the “inlet process”), with a screen, an aerated grit chamber, and solid-liquid separation, all of which are uncovered and exposed to the indoor environment. The secondary treatment is based on activated sludge (hereafter referred to as the “aeration process”), and aeration

is performed in opened aeration tanks with fine bubble diffusers. The wastewater finally moves to an indoor post-treatment process (hereafter referred to as the “outlet process”) with ozone oxidation, coagulation, and filtration. All processes except for the secondary treatment were performed inside buildings. Bioaerosol samples were independently collected at each processing site in the plant to provide insight into any differences in bioaerosol generation, and to compare indoor and outdoor environments in the plant.

The sampling locations were selected according to wind directions and access depending on the operational activities taking place (see the sampling locations in Fig. 1). Indoor samples were usually taken at a distance of 1 to 2 m from the potential sources. Outdoor samples were taken 2 to 4 m downwind from the operations. In all cases, a background sample was taken at an upwind location unaffected by plant operations, usually 10 to 20 m upwind.

2.2. Air Sampling and Analysis

Site visits for sampling were undertaken once a month from October 2009 to September 2010. Microbial investigations were carried out during ordinary work time (1:00 PM to 4:00 PM) at a height of about 1.5 m above the floor of the site being investigated.

Cultivation-based detection in selective media after impaction was applied to detect and characterize the viable components of bioaerosol. Mesophilic heterotrophic bacteria were chosen to be analyzed, because they are prevalent in the environment [9]. Air samples of airborne bacteria and fungi were collected using two air samplers; a single-stage impactor sampler (Buck Bio-Culture Pump model B30120; A. P. Buck Inc., Orlando, FL, USA), and a six-stage viable particulate cascade impactor (Model 10-800; Andersen Inc., Smyrna, GA, USA). The cascade impactor was selected to identify the distribution characteristics of the particular bioaerosol by the aerodynamic particle size, for which the ranges in stages 1 to 6 are >7.0 μm, 4.7–7.0 μm, 3.3–4.7 μm, 2.1–3.3 μm, 1.1–2.1 μm, and 0.65–1.1 μm, respectively. The sum of the colonies counted between stages 3 and 6 corresponds to the respirable fraction. The impaction flow rate for both samplers was set to 28.3 L/min. The duration of air sampling was initially set to 5 min, and then varied according to the environmental conditions at the measurement locations. Bacteria and fungi samples were sampled in sequence.

Trypticase Soy Agar (TSA) (Lot 3087230; BD, Franklin Lakes, NJ, USA), with 500 mg of cycloheximide added to suppress fungal growth, was used as the bacterial culture medium. Airborne fungi were cultured in Sabourand Dextrose Agar (SDA) (Lot 5111476; BD) with 100 mg of chloramphenicol to inhibit the proliferation of bacteria. The culture media were immediately transported to the laboratory after every collection cycle, and cultured in an incubator for 1–2 days at 37°C for bacteria, and for 3–5 days at 25°C for fungi. The colonies were counted, and the measurement was expressed as CFU/m³.

Environmental factors were also measured to analyze any correlation with bioaerosol generation. Temperature and relative humidity were measured with a digital thermohygrometer (Model 608 H-1; Testo, Lenzkirch, Germany). Odor intensity was analyzed with a handheld odor meter (Model OMX-SR; Shinyei, Kobe, Japan). Airborne particulate matter (TSP, PM10, PM2.5, PM1.0) was analyzed using a portable dust monitor (Dustmate; TurnKey Instruments Ltd., Cheshire, UK).

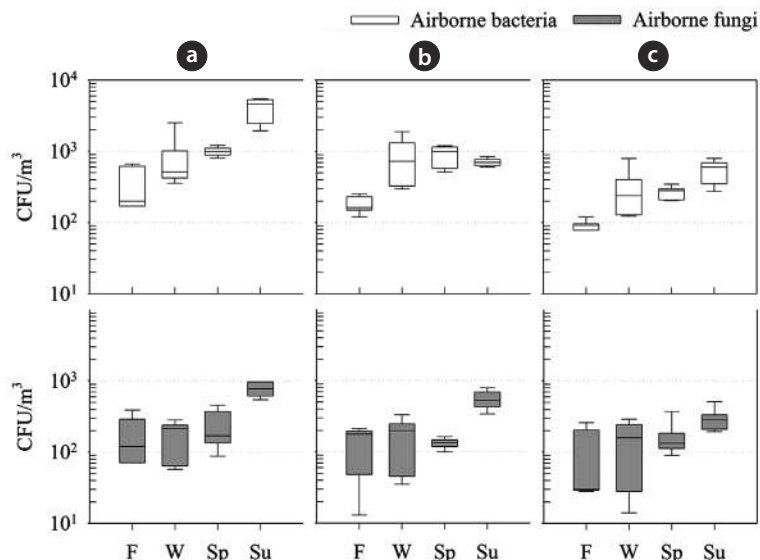


Fig. 2. Seasonal variation in bioaerosol concentrations at the inlet process (a), the aeration process (b), and the outlet process in the wastewater treatment plant (c). The box plots show geometric means and geometric standard deviations. Number of samples for season is 9 each sampling site. F: fall, W: winter, G: spring, S: summer.

2.3. Data Analysis

The microbial contamination was analyzed by season and separately for each step of the treatment process. The statistical significance of the correlation between environmental factors and airborne microorganisms was verified by Pearson and Spearman correlation analysis tests using the SAS/Stat 9.1 (SAS Institute Inc., Cary, NC, USA). The analysis of variance (ANOVA) analysis method was used to verify the statistical significance of the differences between the values of measurements taken for two indoor processes and an outdoor process.

3. Results and Discussion

3.1. Total Culturable Bacteria and Fungi in Air Samples

Culturable bacteria and fungi were detected in all air samples collected at all sites at the scheduled times. The results are summarized in Fig. 2, which presents seasonal variations in bioaerosol concentrations for each step of the treatment process. Concentrations of airborne culturable bacteria generally ranged from 8×10^1 to 5×10^3 CFU/m³, and those of culturable fungi ranged from 1×10^1 to 9×10^2 CFU/m³. In the previous study, the levels of bioaerosol released from twelve swine farms were slightly higher than those of our observation [3]. However, as demonstrated in that study, the difference in overall bioaerosol concentrations was not statistically different.

In municipal waste treatment plants, site workers should not be exposed to levels higher than 5×10^3 CFU/m³ over an 8-hr working day [15], and the threshold limit in occupational atmospheres is 10^4 CFU/m³ for the total bacteria concentration [16]. The observed levels in our study can therefore pose occupational biorisks in an occupational atmosphere.

Fig. 2 shows that the bacterial concentrations at the inlet process were the highest among all sites (ANOVA, $p < 0.01$), as

were the fungal levels at the inlet process (ANOVA, $p < 0.01$). It has been shown in other studies that the pretreatment process and primary clarifier generate the highest concentrations of airborne microorganisms, including bacteria and fungi, among the treatment stages of the wastewater treatment plants [2, 7-9]. The aeration chamber has been recognized as presenting the highest risk for exposure to biological aerosol because of the amount of microbiological pollution generated [11]. Hence, the observed results indicate that potential bioaerosol sources, such as the abundant microorganisms in the raw wastewater, may be subject to mechanical agitation during pretreatment, such as in the grit removal tank, aeration basin, and primary clarifier.

Fig. 2 also shows the seasonal variations in culturable microorganisms in air samples. The maximum exposure was found to occur during the summer at the following mean levels: $4,141 \pm 1,386$ CFU/m³ for culturable airborne bacteria and 765 ± 171 CFU/m³ for culturable airborne fungi, both of which were observed at the inlet process. For indoor sites (inlet and outlet processes), bacterial and fungal levels in air samples were observed in the following descending order: summer > spring > winter > fall, while for the outdoor site, the bacterial level in the summer was not the highest. Karra and Katsivela [9] observed that the heterotrophic bacterial concentration in air decreased during a summer day in the sunshine, accompanied by increasing solar radiation and temperature, while fungal concentration increased slightly. High temperature and solar radiation in the summer could affect the survival of the airborne microorganisms or deactivate them. Correspondence of the bioaerosol level to seasonal changes was significant (ANOVA, $p < 0.01$) for all sampling sites. The seasonal variation in bioaerosols with meteorological conditions such as temperature, humidity, wind speed and direction, solar radiation, and pollutants, has been investigated in previous studies, where it was speculated that these factors should affect the generation levels of bioaerosols outdoors rather than indoors [2, 9]. The relationship between bioaerosol generation and environmental conditions will be discussed in Section 3.3.

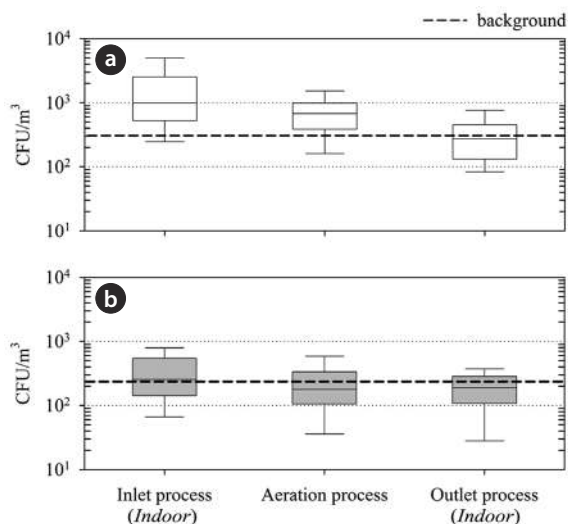


Fig. 3. Summary of bioaerosol concentrations during the field study. The box plots show medians, 5th and 95th percentiles, and outliers. The dotted line indicates background levels of bioaerosols (300 CFU/m³ for airborne bacteria; 220 CFU/m³ for airborne fungi; both are average values over the period of study) at each location. The background data were obtained at a distance of 10 to 20 m from the plant and against the wind direction.

Fig. 3 presents the overall bioaerosol concentrations at the sampling sites during 1 yr of study. A gradual decrease in the concentrations of bacteria and fungi in the air samples was observed after an early step of the treatment process (i.e., inlet process). The mean concentrations of culturable bacteria in air samples were $1,753 \pm 1,693$ CFU/m³ at the inlet process, 735 ± 461 CFU/m³ at the aeration process, and 327 ± 234 CFU/m³ at the outlet process. In comparison, the mean concentrations of culturable fungi were 345 ± 264 CFU/m³ at the inlet process, 238 ± 203 CFU/m³ at the aeration process, and 199 ± 128 CFU/m³ at the outlet process. In total, the concentration of culturable bacteria in the air samples decreased by more than 80% after post-treatment, but those of culturable fungi did so only by around 40%, regardless of seasonal variation. The culturable bacterial concentration in the air samples was, at least, 5 times the culturable fungal concentration at the inlet process; further, the culturable bacterial concentration in the air samples was 3 times that at the aeration process and 1.6 times that at the outlet process. It is important to note that the background levels of bioaerosols were greater than those observed at the outlet process, although far less than those at the inlet process.

The exact microbial concentration cannot be determined using cultivation-based detection due to the fact that there is a methodological limitation of culturing all the individual microorganism colonies existing in an environmental sample [9, 17], leading to underestimation of the airborne living microorganisms as well as underestimation of the risks from toxic or immunopathogenic effects on the workers at a site. Li et al. [10] reported that by polymerase chain reaction and denaturing gradient gel electrophoresis (PCR-DGGE) analysis, pathogenic species such as *Moraxella nonliquefaciens* and *Flavobacterium odoratum*, were detected from the bioaerosol at a municipal wastewater treatment plant. Pascual et al. [7] found *Pseudomonas aeruginosa*

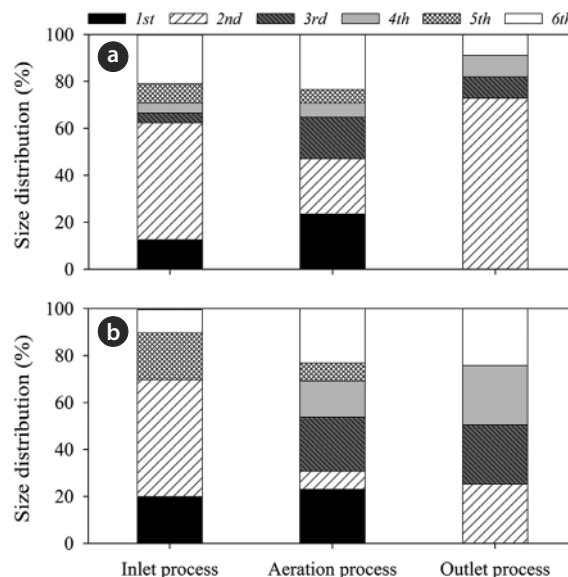


Fig. 4. Size distributions of airborne bacteria (a) and airborne fungi (b) at the sampling locations at the wastewater treatment plant. The aerodynamic diameter ranges for the viable particle sizing sampler were >7.0 μm (stage 1), $4.7\text{--}7.0$ μm (stage 2), $3.3\text{--}4.7$ μm (stage 3), $2.1\text{--}3.3$ μm (stage 4), $1.1\text{--}2.1$ μm (stage 5), and $0.65\text{--}1.1$ μm (stage 6).

at pretreatment and primary clarifiers at a municipal wastewater treatment plant by PCR methods. It is therefore necessary to conduct a qualitative evaluation of airborne microorganisms by identification, which would provide information about contributions from internal generation.

3.2. Distribution of Microorganisms by Bioaerosol Particle Size

Fig. 4 shows a diagram of the distribution of airborne bacteria and fungi according to bioaerosol size at each step of the wastewater treatment plant. At the inlet process, the highest distribution appears in stage 2 (85 CFU/m³ for culturable bacteria and 35 CFU/m³ for culturable fungi), with bioaerosol size is in the range of $4.7\text{--}7$ μm . The lowest distributions were in stages 3 and 4 (7 CFU/m³ for culturable bacteria in both stages and culturable fungi were undetectable). The bioaerosol sizes are in the ranges $3.3\text{--}4.7$ μm and $2.1\text{--}3.3$ μm , respectively. At the outlet process, the highest distribution appeared in stage 2 (57 CFU/m³ for culturable bacteria), where bioaerosol size is in the range of $4.7\text{--}7$ μm , but differences in the distributions of other size ranges were not significant. At this site, culturable fungi show an even distribution in stages 2, 3, 4, and 6. At the inlet process, the ratios of respirable concentration to total concentration (corresponding to bioaerosol sizes of 0.65 to 4.7 μm) were 0.37 and 0.30 for culturable bacteria and fungi, 0.54 and 0.69 at the aeration process, and 0.27 and 0.75 at the outlet process, respectively. In a full-scale composting facility, size distributions of the total airborne microorganisms were evenly distributed in the six stages, and the concentrations at each stage were over 10^4 CFU/m³ [18]. In feedstuff-manufacturing factories, over 70% of airborne bacteria were concentrated in respirable air and more than 60% of airborne fungi consisted of respirable airborne fungi [17]. It has

Table 1. Correlation coefficients^a for geometric means of bioaerosol concentrations compared with environmental parameters

	No. ^b	Airborne bacteria				Airborne fungi			
		Pearson correlation		Spearman correlation		Pearson correlation		Spearman correlation	
		<i>r</i>	<i>p</i> -value	<i>r</i>	<i>p</i> -value	<i>r</i>	<i>p</i> -value	<i>r</i>	<i>p</i> -value
Temperature	24	0.28	<0.01	0.26	0.22	0.78	<0.01	0.90	<0.01
Relative humidity	24	<u>0.41</u>	<0.01	0.21	0.33	<u>0.44</u>	<0.01	<u>0.45</u>	0.03
TSP	27	0.18	<0.01	0.02	0.94	0.04	<0.01	0.05	0.79
PM10	27	0.10	<0.01	0.08	0.69	0.01	<0.01	0.08	0.69
PM2.5	27	0.04	<0.01	0.19	0.34	0.09	<0.01	0.20	0.33
PM1.0	27	0.06	<0.01	0.38	0.05	0.10	<0.01	0.27	0.17
Odor intensity	11	0.78	0.18	0.64	0.05	0.81	<0.01	<u>0.53</u>	0.12

RH: relative humidity, TSP: total suspended particle.

^aFor Pearson correlation coefficient (*r*) and Spearman correlation coefficient (*r*), bold-face: strong association ($r \geq 0.6$); underlined: moderate association ($0.4 \leq r < 0.6$). For *p*-value, significantly correlated ($p < 0.01$); marginally significantly correlated ($0.01 \leq p < 0.05$).

^bNumber of paired data set: non-paired data sets were removed for correlation analysis.

been reported by others [19, 20] that bioaerosols with aerodynamic diameters of 5 μm have greater effect on the alveolus than those with larger diameters, and result in allergic reactions and other serious illnesses. In this study, the outdoor aeration process showed the highest ratio of respirable to total bioaerosol concentration. It is speculated that the secondary generation of microorganisms due to a biological mechanism in the aeration process could contribute to high levels of respirable bioaerosols detected.

Although these bioaerosol distributions are affected by environmental factors, meteorological conditions, and generation sources, the results observed in this study may be site specific. The ratio of respirable bioaerosol was significant, and this poses a potential biological risk, which could be hazardous to workers.

3.3. Effect of Environmental Factors on Bioaerosol Generation

The degree of correlation between airborne microorganisms and environmental conditions at the test sites is summarized in Table 1, which was obtained by applying Pearson correlation analysis and Spearman correlation analysis. Among the environmental factors investigated, air temperature showed a statistically significant correspondence with airborne fungi ($p < 0.01$) for both Pearson and Spearman correlation analyses. In general, Pearson correction is most appropriate for data with a linear relationship, while Spearman correlation is more appropriate for non-linear cases. Hence, Spearman correlation is quite suitable for biological parameters. For the Spearman correlation analysis, the correlation coefficient for temperature was found to be 0.90, while that for relative humidity was 0.45. In our previous study, the correlation between relative humidity and airborne fungi was much more significant than temperature in feedstuff-manufacturing factories [17]. In a sewage treatment plant, both air temperature and humidity were strongly correlated with airborne fungi according to the statistical estimation by Spearman correlation [11]. The growth of airborne fungi is maximized when the relative humidity is greater than 70% [21]. In our field data, the temperatures measured varied from 2.3°C to 33.0°C, and relative humidity varied from 25% to 68%. The relative humidity fluctuated less due to the sampling site being water based.

Fungi are especially versatile, and can utilize different substrates for their growth, and temperature and water availability are the most important factors for fungal growth [22]. For odor intensity, significant associations were found with the generation of both airborne bacteria and airborne fungi. Even odor intensity is not meaningful as an environmental factor in biological development, but it may be a useful indicator of airborne biological issues.

There have been no similar reports on the distributions of airborne microorganisms in swine wastewater treatment plants. However, compared with other studies on municipal wastewater treatment plants [2, 7, 9, 23] and other working sites [16-18, 24], the distribution of airborne microorganisms was found to have a similar pattern. There is no official standard for the exposure to bioaerosols in the indoor area of the studied site. The American Conference of Government Industrial Hygienists (ACGIH) recommends indoor concentrations of bacteria and fungi to be less than 1,000 CFU/m³ [25]. From the results of this study, the indoor pre-treatment process should be of concern to workers, and thus, it is imperative that appropriate institutional standards and maintenance guidance be made available as soon as possible.

4. Conclusions

The concentrations of bioaerosol (mesophilic heterotrophic bacteria and fungi) emitted from the swine wastewater treatment plant varied depending on the step of the process being considered, and by season. The pretreatment process was found to be the major source of bioaerosol. For the aeration process, a portion of the respirable bioaerosol concentration was significant. Our study is the first report on the size distribution of bacteria and fungi aerosolized from a swine wastewater treatment plant using a solid impact air sampler. A significant correspondence between temperature and airborne fungi bioactivity was found, and the odor intensity could be used for detecting high levels of both airborne bacteria and fungi. Our field data should be taken into consideration in the design and maintenance of the swine wastewater treatment plants with respect to occupational safety and health.

Acknowledgments

This study was supported by the Basic Science Research Program through the National Research Foundation of Korea (NRF), funded by the Ministry of Education, Science, and Technology (No. 2009-0088397) and by the Korea Ministry of Environment and human resource development project for energy from waste recycling.

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