Microbial Diversity in the Sawmill Environment: Implications on the Health of Sawmill Workers and Merchants, Nigeria

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Cover Page Footnote
We are appreciative of the laboratory staff of the microbiology laboratory of igbinedion university okada

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MICROBIAL DIVERSITY IN THE SAWMILL ENVIRONMENT: IMPLICATIONS ON THE HEALTH OF SAWMILL WORKERS AND MERCHANTS, NIGERIA

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ABSTRACT

There is an increasingly high suspicion of occupational health diseases amongst workers in sawmill factories. Hence this study aimed to determine the microbial diversity in some sawmill factories in Okada, Edo State, Nigeria to evaluate potential implications of the factory’s processing area on the health of the workers and wood merchants. Bacteria and fungi counts in the bioaerosols within the processing area of each factory was performed with the passive air sampling technique. While in the sawdust samples, microbial counts were carried out with the pour plate technique. Identification of the microbes was performed with macroscopic and microscopic examinations as well as standard phenotypic tests. Mean total viable count (TVC) and total fungi count (TFC) of bioaerosols in the air within the processing area of the sawmill factories were found to be greater than 4162.99 CFU/m³ while mean total coliform count (TCC) was reported as 756.28 CFU/m³. Fisher (F) one-way ANOVA test of the TVC and TCC dataset of bioaerosols from the four sawmill factories indicated no significant difference (p = 0.77 and 0.83 for TVC and TFC respectively) in the mean TVC and TFC of the bioaerosols. Bacterial (Micrococcus, Staphylococcus, Bacillus, Klebsiella and Serratia) and fungal (Saccharomyces, Aspergillus, Cladosporium and Penicillium) species were isolated from the bioaerosols and sawdust samples. Results of this study indicated that workers in the sawmill factories and other wood merchants who visit sawmill processing area are most likely exposed to airborne contaminants that may cause occupational diseases such as ophthalmic irritations and dermatitis. Hence it is recommended that sawmill workers and other visitors wear personal protective gadgets in the processing area of the sawmill factories.

Keywords: Occupational health, sawmill, bioaerosol, sawdust, microbes.

INTRODUCTION

Sawmill factories generate large quantities of organic dust (Demers et al., 2000; Straumfors et al., 2018) that easily disperses into air and has been found to pose a high risk of occupational diseases such as asthma, frequent headaches, chronic bronchitis, phlegm, nasal symptoms and ophthalmic irritations (Demers et al., 1997; Alwis et al., 1999; Idu et al., 2019). The airborne dust is often associated with microorganisms, referred to as bioaerosols (Sawane and Sawane, 2017), which may cause a myriad of adverse health effects on the workers and wood merchants. Several studies have shown that bacteria and fungi are the main airborne microorganisms that are associated with sawdust. Some microbial contaminants of sawdust that have been reported include Micrococcus, Staphylococcus, Klebsiella, Aerococcus, Moraxella, Sphingmonas, Bacillus, Enterobacter, Aspergillus, Cladosporium and Penicillium (Oppliger et al., 2005; Park et al., 2010; Baranu and Edmund, 2019; Idu et al., 2019). This is because before processing the wood, it is stored outside the factory premises thereby creating a favorable environment for the
propagation of these microorganisms (Oppliger et al., 2005). Occupational diseases have increasingly become a source of concern in the world. For instance, in South Korea, the third most common source of occupational diseases is derived from bioaerosols generated from sawmill factories and other workplaces (Park et al., 2010). Occupational health problems associated with bioaerosols generated from sawdust in sawmill factories situated in Switzerland have also been reported (Oppliger et al., 2005). Idu et al. (2019) also indicated occupational health problems in a Nigerian sawmill factory. In order to prevent occupational health problems associated with the sawmill workplace, there is a need to regularly establish occupational exposure levels of the bioaerosols and have a knowledge of airborne contaminants in sawmills. Hence, this study was carried out to determine the microbial diversity in some sawmill factories in Okada, Edo State, Nigeria with a view to evaluate potential implications of the factory’s processing area on the health of the workers and wood merchants.

MATERIALS AND METHOD

Experimental Design

Four sawmill factories situated in Okada, Ovia North-East local government area, Edo State, South-Southern Nigeria were visited between the months of May to July 2020 for sample collection. Each sawmill factory was visited twice weekly for sample collection. In each of the sawmill factories, air and sawdust samples within the vicinity of the sawmill factories were collected. Overall, 24 sawdust samples and 24 bioaerosol samples were collected from all the four sawmill factories examined. One-way analysis of variance of the sample sizes was carried out with the ‘completely randomized design’ technique. Counts of total viable bacteria (TVC), total coliform bacteria (TCC) and total fungi (TFC) was performed on the samples.

Microbial Analysis of Bioaerosol Samples

Bacteria and fungi count in the bioaerosols within the processing area of each factory was performed with the passive air sampling technique or the settle plate method (Pasquarella et al, 2000). Sterile duplicate nutrient agar (NA), MacConkey agar (MA) and Sabourad’s dextrose agar (SDA) Petri dishes held at a height of one meter above the floor of the processing area and one meter away from any obstacle in the processing area were exposed to air for one hour to enable organic dust that is suspended in the air to settle on the Petri dishes. Each exposed Petri dish had a diameter of 8.5 cm with an equivalent area of 60.053 cm². After exposure, the NA and MA Petri dishes were then incubated at 37°C for 24 hours, while the SDA Petri dishes were incubated at room temperature for 72 hours. Colony counts (TVC, TCC and TFC) on the Petri plates were respectively expressed as colony-forming units per cubic meter (CFU/m³) according to the equation stated below (Fekadu and Getachewu, 2015).

\[ N = 5a \cdot 10^4 (bt)^{-1} \]  

\( N \): microbial CFU/m³ of indoor air, \( a \): number of colonies per Petri plate, \( b \): dish surface (cm²), \( t \): exposure time (minutes).

Microbial Analysis of Sawdust Samples

Isolation of bacteria and fungi in the sawdust samples collected from the processing area was carried out with the pour plate technique (Public Health England, 2014). Serial dilution of samples was made up to \(10^{-5}\), with the first dilution made by mixing 25 g of sawdust sample with 225 ml of sterile 1.5% peptone water in a sterile tube. One milliliter of each serially-diluted sample was separately poured into three different duplicate sterile
petri dishes and then respectively mixed with 19 ml of sterile NA, MA and SDA media. The NA and MA Petri dishes were then incubated at 37°C for 24 hours while the SDA Petri dishes were incubated at room temperature for 72 hours. After incubation, microbial colonies were counted with the colony counter and counts on NA, MA and SDA Petri plates, which were expressed as colony-forming units per gram (CFU/g) and were respectively reported as TVC, TCC and TFC.

**Identification of Microbial Colonies**

Identification of the bacterial colonies was performed with standard phenotypic tests (Barrow and Feltham, 2003). The phenotypic tests that were performed on the bacterial colonies included Gram staining, coagulase, catalase, citrate, oxidase, indole, methyl red and Voges Proskauer tests. Identification of fungi was mainly done by macroscopic and microscopic examination using the ‘manual of medically important fungi’ as a guide (Larone, 1995).

**Parasitological Examination**

Sawdust samples were observed for the presence of protozoan parasites with the zinc sulphate centrifugal floatation method as previously prescribed by Parameshwarappa et al. (2012).

**Statistical Analysis**

The NCSS (version 12) data analysis software was used to carry out descriptive statistics of TVC, TCC and TFC. Shapiro–Wilk normality test, Levene test of homogeneity, Fisher (F) one-way ANOVA test for normally distributed datasets with equal variances and Kruskal–Wallis nonparametric one-way ANOVA tests were also performed with NCSS. The test of the hypothesis was considered statistically significant if the achieved level of significance (p) was less than 0.05.

**RESULTS**

**Microbial Counts in Bioaerosols and Sawdust Samples**

Table 1 represents the counts of bacteria and fungi found in the processing area of sawmill factories situated in Okada, Ovia North-East LGA, South-Southern Nigeria. Mean TVC and TFC of bioaerosols in the air within processing area of the factories were found to be greater than 4162.99 CFU/m³; while in the sawdust, mean TVC was recorded as 215000 CFU/g and mean TFC as 92500 CFU/g. Mean TCC of the bioaerosols was reported as 756.28 CFU/m³ while mean TCC of sawdust samples was 215000 CFU/g. The mean TVC and TFC datasets of bioaerosols obtained from all the sawmill factories were normally distributed (p = 0.94 and 0.95 for TVC and TFC respectively) with equal variance (p = 0.62 and 0.68 for TVC and TFC respectively). Fisher (F) one-way ANOVA test of the TVC and TCC dataset of bioaerosols indicated no significant difference (p = 0.77 and 0.83 for TVC and TFC respectively) in the mean TVC and TCC of the bioaerosols. The mean TCC dataset of bioaerosols was non-normally distributed (p = 0.02) with equal variance (p = 0.07). Kruskal-Wallis ANOVA test of the TCC dataset indicated a significant difference (p = 0.04) in the median TCC dataset of aerosols. Fisher (F) one-way ANOVA test also showed that the mean TVC, TCC and TFC datasets of sawdust obtained from all the sawmill factories showed no significant difference (p = 0.63, 0.51 and 0.73 for TVC, TCC and TFC respectively).
Table 1: Microbial counts obtained from sawmill environment

<table>
<thead>
<tr>
<th>Bioaerosols in the processing area of all sawmill factories</th>
<th>Sawdust in the processing area of all sawmill factories</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mean TVC  (CFU/m³)</td>
<td>Mean TVC  (CFU/g)</td>
</tr>
<tr>
<td>N = 24</td>
<td>N = 24</td>
</tr>
<tr>
<td>Mean TCC  (CFU/m³)</td>
<td>Mean TCC  (CFU/g)</td>
</tr>
<tr>
<td>N = 24</td>
<td>N = 24</td>
</tr>
<tr>
<td>Mean TFC  (CFU/m³)</td>
<td>Mean TFC  (× 10⁶ CFU/g)</td>
</tr>
<tr>
<td>N = 24</td>
<td>N = 24</td>
</tr>
<tr>
<td>&gt; 4162.99 ± 0.00</td>
<td>3.37 ± 0.22</td>
</tr>
<tr>
<td>756.28 ± 266.47</td>
<td>2.15 ± 0.80</td>
</tr>
<tr>
<td>&gt; 4162.99 ± 0.00</td>
<td>9.25 ± 2.56</td>
</tr>
</tbody>
</table>

TVC: total viable count; TCC: total coliform count; TFC: total fungi count. Mean count represented as mean ± standard error of mean. N is the total number of samples examined.

Identity of Microbes in Bioaerosols and Sawdust Samples

The microbial community in the bioaerosols and sawdust samples present in the sawmill factories include bacteria, fungi and parasites. The bacteria that were identified in the bioaerosols included *Micrococcus*, *Staphylococcus*, *Bacillus*, *Klebsiella* and *Serratia* species, while the fungi identified were *Saccharomyces*, *Aspergillus*, *Cladosporium* and *Penicillium* species. *Entamoeba* species was the main parasite found in sawdust. The main bacteria and fungi that were present in the air were similar to those that were isolated from the sawdust indicating that the airborne microbes may most likely originate from the wood that was processed in the sawmill factories.
Table 2: Phenotypic characterization of microbial isolates in sawmill environment

<table>
<thead>
<tr>
<th>Samples</th>
<th>Representative isolates</th>
<th>Morphology on Petri plates</th>
<th>Microscopy</th>
<th>Biochemical characteristics of bacterial isolates</th>
<th>Identified microbes</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td>Bacteria</td>
<td>Fungi</td>
<td></td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>1</td>
<td>Macoid colony on NA plate</td>
<td>Macoid colony on NA plate</td>
<td>Gram positive cocci</td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Gram positive cocci</td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td>2</td>
<td>Macoid colony on NA plate</td>
<td>Macoid colony on NA plate</td>
<td>Gram positive cocci in clusters</td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>Gram positive cocci in clusters</td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td>3</td>
<td>Dry colony on NA plate</td>
<td>Red colony on NA plate</td>
<td>Gram positive rods</td>
<td>NP</td>
<td>+</td>
</tr>
<tr>
<td></td>
<td>Red colony on NA plate</td>
<td>Red colony on NA plate</td>
<td>Gram negative rods</td>
<td>NP</td>
<td>+</td>
</tr>
<tr>
<td>4</td>
<td>Macoid colony on NA plate</td>
<td>Macoid colony on NA plate</td>
<td>Gram positive rods</td>
<td>NP</td>
<td>+</td>
</tr>
<tr>
<td></td>
<td>Macoid colony on NA plate</td>
<td>Macoid colony on NA plate</td>
<td>Gram positive rods</td>
<td>NP</td>
<td>+</td>
</tr>
<tr>
<td>5</td>
<td>White colony on SDA plate</td>
<td>White colony on SDA plate</td>
<td>Gram positive buds</td>
<td>Unicellular oval cells that appeared purple upon Gram staining with some cells exhibiting budding</td>
<td>NF</td>
</tr>
<tr>
<td>6</td>
<td>Greenish colonies on the surface of SDA plate</td>
<td>Multiple filaments arising from a single conidiophore carrying chains of conidia</td>
<td>-</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>7</td>
<td>Greenish colonies on the surface of SDA plate</td>
<td>Multiple filaments arising from a single conidiophore carrying chains of conidia</td>
<td>-</td>
<td>+</td>
<td>-</td>
</tr>
<tr>
<td>8</td>
<td>Black colonies on the surface of SDA plate</td>
<td>Black colonies on the surface of SDA plate</td>
<td>Conidiophores arising from septate hyphae with central vesicles that were completely filled with conidia</td>
<td>NF</td>
<td>NP</td>
</tr>
<tr>
<td>9</td>
<td>Greenish colonies on the surface of SDA plate</td>
<td>Greenish colonies on the surface of SDA plate</td>
<td>Multiple filaments arising from a single conidiophore carrying chains of conidia</td>
<td>NF</td>
<td>NP</td>
</tr>
<tr>
<td>10</td>
<td>Greenish colonies on the surface of SDA plate</td>
<td>Greenish colonies on the surface of SDA plate</td>
<td>Saprosporangium spp.</td>
<td>NF</td>
<td>NP</td>
</tr>
</tbody>
</table>

Co, Ca, Ox, Ci, I, Mr and Vp indicate coagulase, catalase, oxidase, citrate, indole, methyl red and Voges Proskauer tests respectively. NP means that the test was not performed. +: positive test. -: negative test.
DISCUSSION

The TVC and TFC in the aerosols within the processing area of all the sawmill factories were greater than (> 300 CFU/Petri plate with an equivalent mean TVC and TCC that was > 4162.99 CFU/m³ (Table 1). The count of Gram-negative bacteria as expressed by the mean TCC was estimated at 756.28 CFU/m³. The results agreed with the findings of Alwis et al. (1999) who obtained bioaerosol levels of microbes in the range of $10^2$ – $10^4$ CFU/m³. It was also similar to those of Park et al. (2010) who reported mean bioaerosol levels of $10^3$ CFU/m³ for bacteria and fungi respectively, as well as to the work of Oppliger et al. (2005) who obtained bioaerosol levels in the range of $10^3$ – $10^4$ CFU/m³ for TVC and TFC including a mean value of $10^2$ CFU/m³ for Gram-negative bacteria in Swiss sawmill factories. The microorganisms associated with bioaerosols in the sawmill factories may be largely due to microbes that are propagated in the bark of the trees during growth in the forest and subsequent storage in the sawmill factories before sawing (Oppliger et al., 2005). Significant exposure of workers and merchants to bioaerosols in the sawmill factories often occurs from the release of high concentration of airborne microbes into the air during sawing of the wood. In line with the findings from this study, Idu et al. (2019) also reported a high concentration of bacteria in the range of $10^6$ – $10^7$ CFU/g of sawdust.

The bioaerosol levels obtained in the sawmill factories examined were compared to recommended exposure limits for bioaerosols in indoor air and occupational environments. Results of the present study indicated that the total microbial load (TVC and TFC) exceeded WHO (Nevalainen and Morawaska, 2009) and Korean (Park et al., 2010) recommended limits for indoor air (1000 CFU/m³ and 800 CFU/m³ for WHO and Korea respectively). Findings from this study also indicated the bioaerosol concentrations (Table 1) were above the occupational exposure guidelines of Switzerland (Oppliger et al., 2005) stipulated for airborne fungi (≤ 1000 CFU/m³) and may be above the recommended values for total bacteria count (TVC) (≤ 10000 CFU/m³ for TVC) but were within the recommended limit for Gram-negative bacteria (TCC) (≤ 1000 CFU/m³ for TCC). The high concentration of total viable bacteria and fungi in the air within the processing area of sawmill factories in Okada, Edo State, Nigeria may most likely induce occupational asthma and other respiratory diseases in workers and merchants if adequate precautionary measures are not regularly observed in the work area during sawing of wood. Micrococcus, Staphylococcus, Bacillus, Klebsiella and Serratia species were the bacteria that were isolated from the air and sawdust samples collected from the sawmill factories (Table 2) while the fungi that were found include Saccharomyces, Aspergillus, Cladosporium and Penicillium. The parasite that was detected in sawdust samples was mainly Entamoeba species. Park et al. (2010), Oppliger et al. (2005), Idu et al. (2019) as well as Baranu and Edmund (2019) have all reported similar microorganisms in other sawmill factories. The bacteria in the organic dust may infiltrate into the lungs and release endotoxins that may most likely cause adverse health effects in humans that are exposed to sawdust. The airborne fungi may also cause chronic infections such as pulmonary aspergillosis and aspergilloma, particularly in immune-compromised humans that are exposed to sawdust.

CONCLUSION

High levels of bacteria and fungi were present in bioaerosols of the organic dust generated by processing operations in all the sawmill factories examined as indicated by WHO and other international
occupational exposure limit guidelines. Thus, workers in these sawmill factories and other wood merchants who visit the sawmill processing area are most likely exposed to airborne contaminants that may cause occupational asthma, ophthalmic irritations and other respiratory diseases as well as dermatitis. Hence it is recommended that sawmill workers and other visitors within the processing area wear personal protective gadgets such as nose masks, respirators, helmets and safety boots.

REFERENCES


