

2 GLOBAL WARMING POTENTIALS OF MUNICIPAL SOLID
3 WASTE DUMPSITE IN CALABAR METROPOLIS, CROSS
4 RIVER STATE

5
6 ABSTRACT

7 Total anthropogenic greenhouse gases have continued to increase with municipal
8 solid waste dumpsites contributing immensely to the high concentrations of
9 greenhouse gases with high warming potentials. This study is aimed at
10 determining the global warming potentials of municipal solid waste dumpsite in
11 Calabar metropolis. Soil, leachate and solid wastes samples were collected from
12 the dumpsite and subjected to standard microbiological analysis. The samples
13 were examined for temperature, p^H , mean heterotrophic bacterial/ fungal counts
14 and greenhouse gases emissions from the dumpsite using Combustible Gas Leak
15 Detector. The mean temperature values for both soil and leachate samples ranged
16 from 82⁰F-83⁰F while the mean p^H values ranged from 6.57-7.0. The proportion
17 of Carbondioxide, Methane and Nitrous oxide in the studied dumpsite has
18 significantly increased to 39%, 161% and 19% respectively. The mean total
19 viable aerobic heterotrophic bacterial count in both leachate and soil samples
20 ranged from 1.7×10^3 - 8.0×10^3 cfu/ml and 1.2×10^4 - 8.0×10^4 cfu/g, while the
21 mean total viable fungal counts for both leachate and soil samples ranged from
22 1.0×10^5 – 5.0×10^5 cfu/ml and 2.1×10^3 – 6.0×10^3 cfu/g. the prevalent bacteria
23 isolate from the dumpsite soil, leachate and decomposing solid waste include:
24 *Escherichia coli*, *Bacillus sp*, *Pseudomonas aeruginosa*, *Enterobacter sp*,
25 *Klebsiella sp*, *Proteus sp*, *Salmonella sp*, *Staphylococcus aureus*, *Micrococcus*
26 *luteus*, and *Methanococcus sp*. The fungi isolated include: *Candida tropicalis*,
27 *Aspergillus sp*, *Penicillium sp*, *Candida parapsilosis*, *Candida albicans*, and
28 *Saccharomyces sp*. Statistical analysis of the bacterial and fungal counts showed
29 significant difference ($p < 0.05$) between the various sampling points. This study
30 reveals the health, environmental, and climate hazard that could result from
31 indiscriminate dumping of untreated wastes.

32 **Key words:** Global warming potential, solid waste, Lemna dumpsite, greenhouse
33 gases

34

35

INTRODUCTION

36 The increasing awareness of global warming all over the world,
37 and most recently its association with solid waste biodegradation is a
38 cause for concern as temperature ranges around such waste dumpsites
39 are of increase than normal. Warming of the climate system is
40 unequivocal since the 1950s with observed changes (IPCC, 2014).
41 Most researches on the causes of this high temperature have reported
42 greenhouse gases (GHG) emitted from microbial activities of solid
43 waste dumpsite (Bassey, 2012). Lemna solid waste dumpsite formerly
44 known as Ikot-Effanga Mkpa dumpsite is an open dump that
45 encourages the breeding of rodents and proliferation of most
46 pathogenic bacterial and fungal species with no report of the dangers
47 associated with these emissions (Eja *et al.*, 2010; Bassey *et al.*, 2015).

48 Waste is any substance, solution, mixture for which no direct
49 use is envisaged but which is transported for reprocessing, elimination
50 by incineration or other methods of disposal (Yakowitz, 1988). With
51 urban industrialization, social development and population increase,

52 solid waste production are growing rapidly making pollution a serious
53 problem (Khupe, 1996; Yaliang, 1996).Solid waste disposal poses
54 threat to humans, animals and the environment at large. Like chemical
55 hazards, etiologic agents might be dispersed in the environment
56 through water and wind.

57 Waste management in developing countries is usually equated
58 with land disposal or discharge water (Cilinskis and Zaloksnis, 1996).
59 This method of waste management is unscientific and causes nuisance
60 to the public and most of all results in global warming as a result of
61 GHG emitted into the atmosphere. This gases alter the climate and
62 also deplete the ozone layer when waste is dumped on land, soil
63 micro-organism including fungi and bacteria, readily colonize the
64 waste carrying out the degradation and transformation of degradation
65 (organic) materials in the waste (Stainer *et al.*, 1989). Micro-
66 organisms in the waste dump use the waste constituents as nutrients,
67 thus detoxifying the materials as their digestive processes breakdown
68 complex organic molecules into simple less toxic molecules (Pavoni
69 *et al.*, 1975).

70 Calabar city does not have a sanitary landfill. Improper disposal
71 of untreated municipal solid waste is not only harmful to human's
72 health but also constitute a threat to the ecological environment
73 (Yaliang, 1996). Temperatures around waste dumps seemed to be
74 higher than normal, hence, the need to study keenly the global
75 warming potentials associated with waste biodegradation in Lemna
76 waste dumpsite.

77 **MATERIALS AND METHODS**

78 **The Study Area and Sampling Site**

79 The Lemna dumpsite in Calabar Municipality which is where this
80 study was carried out, lies geographically along longitudes $08^{\circ}18'E$ and
81 $08^{\circ}26'E$ Greenwich meridian and latitudes $04^{\circ}55'N$ and $04^{\circ}58'N$ of
82 the equator. In the North, Calabar Municipality is bounded by
83 Odukpani and Akamkpa Local government Area, at the East by the
84 Great Kwa River. At the south, it is bounded by the Calabar River and
85 Calabar South Local government Area. It has an area of
86 331.551 square kilometres. Calabar municipality is characterized by a
87 double maxima rainfall regime which occurs in June and September.

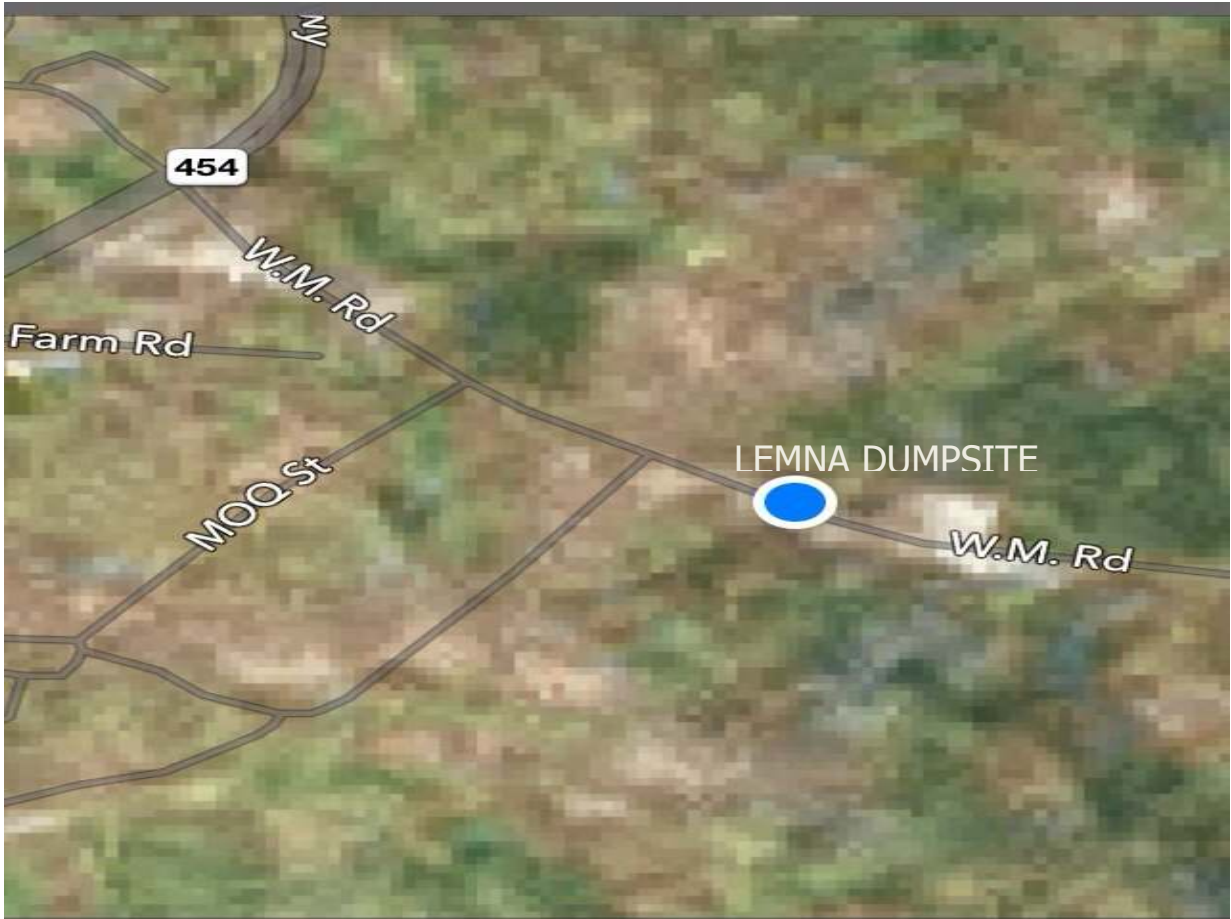
88 The annual temperature is 28⁰cwith a high evapotranspiration and an
89 average humidity of 90%.

90 The vegetation of the study area is characterized by mangrove
91 swamp and rainforest, but due to human activities like cutting down of
92 tress, for roads, building of houses, schools, petrol stations, hotels and
93 market it has resulted in the depletion of the rainforest.

94 It has an annual rainfall of 3000mm and a harmattan wind blowing
95 over the area in December and January respectively.

96 The sampling site was a dump located within Lemma road
97 Calabar Municipality. It is a waste dump site used majorly by the
98 Calabar urban development authority. The dump site was measured
99 with tape and mater rule. The length of the landfill was 960m, the
100 width 430m. Sampling station was established on the waste dumpsite
101 and was represented as soil (S), leachate (L), solid waste (SW) and air
102 sample (AS).

103



104

Figure 1: Map of the study area



107 **Figure 2: Lemna road waste dumpsite**

110

111 **Sample Collection**

112 **Collection of Soil Sample**

113 For soil sample collection, the wastes were first removed using
114 a garden rake to expose the soil under the waste dump from where the
115 soil samples were collected. The soil samples were taken at about
116 15cm depth with the use of hand-driven auger and immediately
117 transported to the laboratory in labeled polythene bags in ice-cold
118 boxes at approximately 4°c for microbiological analysis. A control
119 sample was taken from a location of about 500 meters away from the
120 dumpsite.

121 **Collection of Decomposing Solid Waste Sample**

122 A sterile large wooden spatula that had been properly rinsed
123 with sterile distilled water was used to collect the decomposing wastes
124 into sterile glass petri- dishes and sealed with masking tape. The
125 samples were immediately transported to the laboratory for
126 microbiological analysis or stored in a refrigerator at 4°c until they
127 were needed for analysis.

128

129 **Collection of Leachate Sample**

130 The method described by Bassey *et al.*, 2015, was employed for
131 leachates sampling. In this method, Pvc pipes were cut into four parts,
132 each of 1m and 0.5m in length. The base end of each pipe was
133 permanently sealed with a pipe cover and an adhesive while the top
134 ends were just fitted with pipe covers. The pipes (both 1m and 0.5m
135 length) were perforated evenly at considerable distances from their
136 base ends to allow for water (leachates) percolation and collection.
137 The perforated pipes were then buried at the studied dumpsite for a
138 period of 3-4 weeks to allow percolation of leachates. After 3-4
139 weeks, sterile enema pumps or sterile syringes were used to collect the
140 leachate into sterile bottles and transported to the laboratory for
141 microbiological and physicochemical analysis.

142 **Preparation of Diluent and Media**

143 A ten-folds serial dilution of the sample was made by weighing one
144 gram of sample into 9mls of sterile distilled water to obtain 10^{-1}
145 dilution, further dilutions were made until 10^{-9} diluent was obtained.

146 The media used to grow bacterial were nutrient Agar, Eosin
147 methylene Blue Agar, *Salmonella-shigella* Agar, while the media used
148 to grow fungi was potato Dextrose Agar. All were originally in a
149 powdered form the media is usually prepared by weighing out a
150 certain gram, depending on the number of plates to be prepared. The
151 weighed quantity is then mixed with the appropriate amount of sterile
152 distilled water in a conical flask. The conical flask is usually plugged
153 with enough cotton wool and then placed in an autoclave for
154 sterilization at 121⁰c for 15mins after which the media is removed
155 from the autoclave and allowed to cool.

156 **Determination of P^H and Temperature**

157 The samples were made in suspension (solution) and p^H and
158 temperature probes were inserted into it. The p^H and temperature was
159 recorded accordingly.

160

161 **Microbiological analysis of soil and decomposing waste sample**

162 One gram(1g) of the sample was dissolved in 9ml sterile
163 distilled water from the solution, ten-fold serial dilutions in the range
164 of 10⁻¹ – 10⁻⁹ were prepared .One milliliter(1ml) aliquot from the

165 dilution of 10^{-4} of each samples were aseptically transferred into
166 freshly prepared nutrient agar plates then eosin methylene blue agar
167 and salmonella-shigella agar as selective media for isolation and
168 identification while 0.1ml from the dilution of 10^{-3} was aseptically
169 transferred into potato dextrose agar plates were incubated at 37°c (24
170 hours) for bacterial isolates and $28 \pm 0.2^{\circ}\text{c}$ (3 to 5 days) for fungal
171 isolates. Visible colonies of between 30-300 were multiplied by the
172 reciprocal of the dilution factors and recorded as colony forming units
173 per gram (cfu/g) of waste.

174 **Microbiological analysis of air at dumpsite**

175 Settle plates method described by Bassey et al., 2015, was
176 adopted for this analysis. According to this method, nutrient agar in
177 triplicate was exposed at the dumpsite (10m, 20m, 30m away from the
178 dump) 1hour between 10-12 noon. The plates were later covered and
179 transported to the laboratory for incubation at $25\text{-}28^{\circ}\text{c}$ for 24 hours. The
180 bacterial colonies that appeared on the plate were counted.

181 **3.6.3 Microbiological analysis of leachate**

182 Leachates from such a dumpsite are often turbid therefore, serial
183 dilutions were made. One milliliter (1ml) of the leachate was dissolved

184 in 9ml of sterile distilled water to give 10^{-1} dilution from which further
185 dilution up to 10^{-9} was made (Asikong and Udofia, 2005). The dilutions
186 were then further analyzed as those of decomposed waste and soil
187 sample previously described above. The colony counts were expressed
188 as colony forming units per milliliter of leachate (cfu/ml).

189 **Isolation, characterization and identification of bacteria in the waste**
190 **dump site**

191 Pure cultures of bacteria were obtained by aseptically streaking
192 representative colonies of different morphology types which appeared on
193 the culture plates onto freshly prepared Nutrient agar plates which were
194 incubated at 28°c for 24hours. Discrete bacteria colonies developed and
195 were sub-cultured on nutrient agar slope, incubated at 28°c for 24hours.
196 The nutrient agar slope served as pure culture stock for subsequent
197 characterization tests. The pure cultures were identified based on their
198 cultural, physiological and morphological characteristics. For tests
199 conducted for characterization and identification, see appendix.

200 **Isolation, characterization, identification of fungi in the waste**
201 **dumpsite**

202 Pure cultures of fungi were obtained by sub-culturing discrete
203 colonies into freshly prepared sabouraud dextrose agar plates and
204 inoculated at 28⁰c for 5-7 days. The fungal isolates which developed
205 were further sub-cultured onto agar slope and incubated at 28⁰c for 5 to
206 7days.The isolates which developed were pure cultures which were
207 stored in the refrigerator as stock cultures for subsequent characteristic
208 test. Macroscopic and microscopic examinations of fungal growth were
209 carried out.

210 **Determination of Greenhouse Gas Emissions From the Dumpsite**

211 The emission of greenhouse gases from the studied dumpsite was done
212 using combustible gas leak detector.

213

214 **RESULTS**

215 The frequency of occurrence of bacterial and fungal species isolated
216 from samples obtained from Lemna waste dumpsites are presented in
217 Tables 1 and 2. The table(s) identifies the bacteria and fungi with the
218 highest and lowest frequency of occurrence. From the results obtained,
219 *Escherichia coli* had the highest frequency of occurrence while
220 *Micrococcus luteus* had the lowest frequency of occurrence Table 1.

221 *Penicillium* were observed as the most occurring fungal species while
222 *Candida tropicalis* had the lowest frequency of occurrence.

223 The mean bacterial and fungal counts of isolates from the different
224 sources sampled are presented in Tables 3 and 4. The results revealed
225 *Escherichia coli* with the highest mean counts of 27 ± 15.60 cfu/g of
226 decomposing solid wastes while *Pseudomonas aeruginosa* had the least
227 mean counts of 5 ± 3.00 cfu/g of decomposing solid wastes (Table 3).
228 Table 4 also identified the fungus with the highest and lowest mean
229 count for the different samples. *Candida albicans* had the highest mean
230 count of 20 ± 11.60 while *Candida tropicalis* had the lowest mean count
231 of 5 ± 3.00 . Table 5 and 6 showed the mean temperature and P^H values
232 for both soil and leachate samples obtained from the dumpsite.

233

234 **Table 1:** Distribution and frequency of occurrence of bacteria species
 235 isolated from the dumpsite samples

| Isolated Bacteria species | Frequency of occurrence (%) of Bacteria from different sources. | | | |
|-------------------------------|---|------------|------------|------------|
| | S | L | SW | AS |
| <i>Escherichia coli</i> | 80(16.50) | 55 (10.83) | 44 (17.0) | 25 (7.50) |
| <i>Bacillus sp</i> | 63(6.19) | 48 (9.45) | 39 (15.1) | 30 (9.00) |
| <i>Pseudomonas aeruginosa</i> | 52(10.72) | 38 (7.48) | 20 (7.72) | 15 (4.50) |
| <i>Enterobacter sp</i> | 37(7.63) | 50 (9.84) | 30 (11.50) | 20 (6.00) |
| <i>Klebsiella sp</i> | 49 (10.10) | 33 (6.50) | 27 (10.42) | 40 (12.00) |
| <i>Proteus sp</i> | 40(8.25) | 44 (8.70) | 15 (5.80) | 38 (11.34) |
| <i>Salmonella sp</i> | 39 (8.04) | 30 (5.91) | 18 (6.94) | 40 (12.00) |
| <i>Staphylococcus aureus</i> | 43 (8.87) | 64 (12.50) | 21 (8.11) | 45(13.43) |
| <i>Micrococcus luteus</i> | 29 (5.98) | 74 (14.40) | - (-) | 56 (16.71) |
| <i>Methanococcus sp</i> | 53 (10.93) | 73 (14.40) | 45 (17.40) | 26 (7.80) |
| Total | 485 | 508 | 259 | 335 |

236

237 **Key:** S = soil, L = leachate, Sw = solid waste , AS = Air sample

238

239

240

241 **TABLE 2:** Distribution and frequency of occurrence of fungal species
 242 isolated from dumpsite samples.

| Isolated Fungal Species | Frequency of occurrence (%) of Fungal from different sources | | | |
|-----------------------------|--|------------|------------|-----------|
| | S | L | SW | AS |
| <i>Candida albicans</i> | 60(22.00) | 45(23.40) | 35(16.00) | 0(0) |
| <i>Aspergillus sp</i> | 40(15.00) | 30(15.62) | 38(17.40) | 25(45.50) |
| <i>Saccharomyces sp</i> | 45(16.50) | 32(16.70) | 58(26.50) | 30(45.60) |
| <i>Candida tropicalis</i> | 20(7.33) | 15(7.80) | 18(8.23) | 0(0) |
| <i>Penicillium sp</i> | 58(21.23) | 39(20.30) | 50(22.83) | 0(0) |
| <i>Candida parapsilosis</i> | 50(18.30) | 31(16.15) | 20(9.13) | 0(0) |
| Total | 273 | 192 | 219 | 55 |

243

244 **TABLE 3:** Mean Count of bacteria species isolated from dumpsite
 245 samples.

| Isolated Bacterial species | Mean count of Bacterial spp from different sources | | | |
|-------------------------------|--|------------|-----------|-----------|
| | S | L | SW | AS |
| <i>Escherichia coli</i> | 27 ±15.6 | 18.3± 10.6 | 14.7±8.50 | 8.3 ± 4.8 |
| <i>Bacillus sp</i> | 21 ± 1.20 | 16 ± 9.30 | 13±8.00 | 10 ± 6.00 |
| <i>Pseudomonas aeruginosa</i> | 17 ± 1.00 | 12.7 ±7.3 | 7 ±4.00 | 5 ± 3.00 |
| <i>Enterobacter sp</i> | 12 ± 7.00 | 17 ± 9.60 | 10±6.00 | 7 ± 4.00 |
| <i>Klebsiella sp</i> | 16.3 ±9.40 | 11 ± 6.40 | 9±5.20 | 13.3±7.70 |
| <i>Proteus sp</i> | 13.3±7.70 | 15 ± 8.50 | 5±3.00 | 12.7±7.30 |

| | | | | |
|------------------------------|------------|------------|----------|------------|
| <i>Salmonella sp</i> | 14.3 ±8.30 | 21.3 ±12.3 | 7 ±4.00 | 15 ± 8.70 |
| <i>Staphylococcus aureus</i> | 13 ± 8.00 | 10 ± 6.00 | 6±3.50 | 13.3±7.70 |
| <i>Micrococcus luteus</i> | 15.6 ±9.00 | 19.6 ±11.4 | 0(0) | 22.0±13.00 |
| <i>Methanococcus sp</i> | 17.7 ± 10 | 24.3 | 15 ±8.70 | 8.6 ± 5.00 |
| | | ±14.00 | | |

246

247 **TABLE 4:** Mean Count of fungal species isolated from dumpsite
248 samples.

| Isolated Fungal Species | Mean Count of Fungal species from different sources | | | |
|-----------------------------|---|------------|-------------|-----------|
| | S | L | SW | AS |
| <i>Candida albicans</i> | 20± 11.60 | 15 ±8.70 | 11.7 ± 6.70 | 0(0) |
| <i>Aspergillus sp</i> | 13.3± 7.70 | 10±5.80 | 12±7.30 | 8.3±4.80 |
| <i>Saccharomyces sp</i> | 15± 8.70 | 10.7±6.20 | 19.3±11.20 | 10±6.00 |
| <i>Candida tropicalis</i> | 6.7± 4.00 | 5±3.00 | 6±3.50 | 0(0) |
| <i>Penicillium sp</i> | 19.3± 11.20 | 13±7.30 | 16.7±9.60 | 0(0) |
| <i>Candida parapsilosis</i> | 17± 9.60 | 10.3± 6.00 | 7± 4.00 | 0(0) |
| Total | 273 | 192 | 219 | 55 |

249

250 **TABLE 5:** Temperature of the samples

| Temperature Reading (Days) | Sample code | | | |
|----------------------------|----------------|----------------|----------------|-----------------|
| | S ₁ | S ₂ | L ₁ | L _{2a} |
| 1 | 82 | 82 | 83 | 82 |

2 82 83 83 82

Total 164 165 166 164

Mean 82 82.5 83 82

251 **Key:** S₁ = soil sample I, S₂ = Soil sample II, L₁ = Leachate sample I,

252 L₂ = Leachate sample II

253 **Note:** Temperature in °F to °C ($T^{\circ F} - 32 \times 5/9$)

254 Temperature in °C to °F ($T^{\circ C} \times 9/5 + 32$)

255 **TABLE 6:** Mean pH values of samples collected from the refuse

256 dump site.

| pH Reading (Days) | SAMPLE CODE | | | |
|------------------------------|----------------------|----------------------|----------------------|----------------------|
| | S₁ | S₂ | L₁ | L₂ |
| 1 | 7.0 | 7.0 | 7.0 | 7.0 |
| 2 | 7.0 | 6.57 | 6.9 | 7.0 |
| Total | 14 | 13.6 | 13.9 | 14.0 |
| Mean | 7.0 | 6.98 | 6.95 | 7.0 |

257

DISCUSSION

258 The gradual increase of temperature around municipal solid waste
259 dumpsites within Calabar Metropolis were glaring over the years with serious
260 impacts to the immediate environment and biodiversity. The increase is
261 attributed to the continuous emissions of greenhouse gases (GHGs) from the
262 municipal solid waste dumpsites (Bassey, 2012; Bassey *et al.*, 2015). The
263 temperature values of soil and leachate samples from this study ranged from
264 82⁰F to 83⁰F. While the mean temperature value for soil samples was 82.3⁰F and
265 83⁰F for leachates. The pH values ranged from 6.6 to 7.0. The mean pH value
266 for soil sample was 6.9 and 7.0 for leachates. The observed temperature ranges
267 in this study are such that support the growth of mesophilic bacteria
268 (mesophiles). From this study, mesophilic bacteria species are observed to be
269 chiefly involved in biodegradation of the organic components of waste at the
270 studied dumpsites (table 1 and table 3). This implies that, the digestion and
271 decomposition of organic matter present in the waste is only achievable with the
272 high presence of these organisms. In the decomposition process of the waste, it
273 is observed that psychrophiles were more prevalent. This again is a clear
274 indication that psychrophiles are the major organisms that actually started the
275 decomposition process as they are active at lower temperature up to 55⁰F thus
276 generating heat in the process. As the temperature of the decomposing waste
277 reaches 50-100⁰F, the psychrophilic bacteria were seen displaced by the

278 mesophilic bacteria which then continue the biodegradation. The microbial
279 activities of the mesophilic bacteria often elevate the temperature of the compost
280 (decomposing solid waste) which now gives way for the thermophilic flora such
281 as *Bacillus*, *Aspergillus* as reported in this study. The degree of acidity (pH),
282 reported in this study for all the stations of the waste dump site ranged from 6.5
283 to 7.0. Hagerty *et al.*, (1993), reported that, the initial pH of solid waste is
284 between pH 5.0 and 7.0 for refuse which is about 3 days old. The pH of the
285 waste drops to 5.0 or less in the parts 2-3 days of composing and then begins to
286 rise to about 8.5 for the remainder of the aerobic process.

287 According to the soil classification by Odu *et al.*, (1985), the degree of
288 acidity for sampled soil and leachate ranged from slightly acidic to neutral. The
289 frequency of occurrence (%) of isolated bacterial for the different samples
290 showed the organisms that occurred most. Among all the bacteria isolated from
291 the different samples only *Micrococcus* was not isolated from solid waste (SW).
292 The frequency of occurrence of isolated fungi for the different samples showed
293 the fungal that occurred most. Among all the fungi isolated from the different
294 samples *Candida albicans*, *Candida tropicalis*, *Penicillium spp*, *Candida*
295 *parapsilopsis* was not isolated from the Air sample.

296 The total aerobic bacterial count was highest in leachate samples and
297 lowest in solid waste (SW) samples. The order of decrease in the bacterial
298 counts for all the samples indicates that, solid waste (259) < Air < soil (485)

299 <leachate (508). The total viable fungal count was highest in the soil samples
300 and lowest in Air samples. The order of decrease in the fungal counts for all the
301 samples however shows that, Air (552) < leachate (192) < solid waste (219) <
302 soil (273). The bacterial counts were higher than the corresponding fungal
303 counts. This is because the temperature range (82⁰f to 83⁰f) supports the
304 proliferation of the bacteria than fungi.

305 This study shows the frequency of occurrence of bacterial and fungal
306 species isolated from Lemna waste dumpsite in connection with their
307 biodegradation potentials of organic wastes that often results to the emission of
308 greenhouse gases. The prevalent bacteria species isolated from the dumpsite
309 include: *Escherichia coli*, *Micrococcus luteus*, *Bacillus* sp, *Proteus* sp,
310 *Pseudomonas aeruginosa*, *enterobacter* sp, *Klebsiella* sp, *Salmonella* sp,
311 *Methanococcus* sp, *Staphylococcus aureus*. Of all the different genera of
312 bacteria isolated from the waste dumpsite, *Escherichia coli* had the highest
313 frequency of occurrence while *pseudomonas aeruginosa* had the lowest
314 frequency of occurrence.

315 Most of the bacterial isolates reported in this study have been reported
316 (Eja et al., 2010; Bassey et al., 2015; Ekundayo, 2001) to be associated with
317 waste and waste biodegradation. *Bacillus* and *Pseudomonas species* were
318 reported by Gray (1967) to be associated with waste. *Bacillus*, *E. coli*, *Klebsiella*
319 and *Pseudomonas* were also reported by cook et al, (1964). Ekundayo, (1977)

320 reported *Bacillus*, *E. coli*, *Proteus*, *Pseudomonas*, *Staphylococcus*,
321 *Streptococcus* among others. *Pseudomonas* has been widely reported to be
322 associated with waste (Sabry, 1992). Fungal species isolated from the Lemna
323 road dumpsite include *Candida albicans*, *Aspergillus spp*, *Saccharomyces sp*,
324 *Candida tropicalis*, *Penicillium spp*, *Candida parapsilosis*. From the results
325 obtained, *Candida tropicalis* recorded the lowest frequency of occurrence while
326 *Saccharomyces sp* recorded the highest frequency of isolation. All fungi isolates
327 reported in this study have been reported to be associated with waste and waste
328 biodegradation (Ekundayo, 1977; Ikpedu, 1980) reported *Aspergillus* and
329 *Saccharomyces*.

330 The study has revealed the presence of various bacteria and fungi known
331 to be associated with waste biodegradation with subsequent release of
332 greenhouse gases that has contributed immensely to the upscale of temperatures
333 within the study area. Amongst these bacterial species, *Methanococcus spp* was
334 the most prevalent organism in sampling periods when methane gas emissions are
335 relatively high. Bassey *et al.*, 2012, reported that the upscale of temperatures
336 around Lemna waste dumpsite was as a result of the high percentage emissions
337 of greenhouse gases with methanogens been the most occurring organisms. The
338 percentages of greenhouse gases (Carbondioxide CO₂, Methane CH₄ and
339 Nitrous oxide N₂O) measured at the studied dumpsites greatly contributed to the
340 potential warming of temperatures around the dumpsites. If the activities of the

341 encountered bacteria, fungi and yeast species are properly harnessed, it can be
342 used in future treatment plant, in Nigeria for improving the bioconversion of
343 waste compost into organic fertilizer for optimum use in agriculture and
344 horticulture. All the bacterial genera reported in this study with exception of
345 *Enterobacter sp* are potential pathogens (Bassey et al., 2015; Cheesbrough,
346 2005; Cook et al., 1967; Ekundayo, 2001). Also, all the fungal genera reported
347 in this study with exception of *Penicilium sp* are potential pathogens. The
348 presence of these potential pathogens (bacterial and fungal) reported in this
349 study may be attributed to the disposal of raw human faecal discharges and other
350 human wastes at the dumpsite that possibly encouraged their proliferation.

351 The indiscriminate dumping of waste around residential areas and other
352 ecologically sensitive area plays an inevitable role in global warming and also
353 pose serious health challenges. In a case where wastes are dumped
354 indiscriminately, microbes set in to carry out decomposition. During the process
355 of decomposition leachate are formed, if close to a river source the leachate
356 formed flows in and contaminate the river which then becomes hazardous to the
357 community nearby. Also, during the process of decomposition greenhouse gases
358 are formed such as methane which contributes to the depletion of the ozone
359 layer. All these consequences are as a result of the indiscriminate dumping of
360 waste that has over the years been under estimated. Therefore, Cross river state
361 waste management authorities must as a matter of urgency direct her efforts

362 towards the treatment of waste before disposal as to minimize the health hazards
363 associated with dumping of waste.

364 **Conclusion**

365 The present study reveals the potential hazards of indiscriminate dumping
366 of solid waste on the environment at large, with regards to global warming
367 contributions from waste biodegradation. Therefore, the results highlight the
368 importance of proper collection, transportation, disposal and treatment of solid
369 waste in an environmentally friendly manner that will not deface the aesthetics
370 of the ecosystem. Calabar with a vast rising population of over 400,000 people
371 has led to high generation of municipal solid waste with decreased space for
372 disposal. The high concentrations of carbon dioxide, methane and nitrous oxides
373 around the studied dumpsite is a wakeup call for more research tailored towards
374 generating necessary measures that can arrest the risk associated with the
375 continual emission of these greenhouse gases (GHGs) from the dumpsites.

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