



Assessment of bacteria and SPM in the indoor air of households of urban area of Jammu (J&K), India

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Abstract: The present study was conducted to assess status of bacteria (Gram +ve and Gram –ve) in the indoor air of Households located in Jammu city. The study area was divided into eight Sites. At each site two Households were selected randomly and in each Household sampling of SPM ($\mu\text{g}/\text{m}^3$) and bacteria (CFU/ m^3) was done twice at three sub sites. Total bacterial count and SPM was found to be maximum (9308.24 CFU/ m^3 and 1006.12 $\mu\text{g}/\text{m}^3$ respectively) in Households near Water Body and total bacterial count and SPM was minimum (5251.00 CFU/ m^3 and 659.09 $\mu\text{g}/\text{m}^3$ respectively) in Households near Hospital. A significant positive correlation (r) was found between SPM and no. of Gram +ve (+0.18 to +0.78) as well as between SPM and no. of Gram –ve (+0.21 to +0.76) bacteria in the study area.

Keywords: Household, Indoor SPM, Bacteria, Correlation

INTRODUCTION

Air pollutants are prime culprits responsible for polluting both indoor as well as outdoor environment. Indoor air quality refers to quality of air in buildings occupied spaces. It should be probably be called indoor air pollution. Our buildings have undergone radical changes over past few decades thereby resulting in less opportunity to exchange indoor air with outdoor air. This has led to concentration of air pollutants like dust, CO_2 , bacteria etc within the building (Purohit and Ranjan, 2005).

The causal agents of illness and stress can be of chemical, physical or biological origin and have a sizeable impact on productivity. Biological contamination of environment has received great attention in recent years as a possible cause of illness at home and at work place. The interrelationships of building materials, construction, services and spatial arrangements with their environment, occupants and contents are complex and can influence our health and comfort in buildings. Microorganisms are always present in outdoor air but their number and type's changes with time of day, weather, season, geographical location and with the presence of local spore sources (Nair *et al.* 1996).

Most people spend 90% of their time indoors. Indoor air quality plays a greater role in the health of people who spend most of their time indoors i.e. the very young, the very elderly and those who are chronically ill. Air inside homes, offices, schools and other buildings can be more polluted than outdoor air because heating; cooling and

ventilation systems are common sources of bacterial as well as fungal growth.

The aim of present study is to assess the status of bacteria and SPM in the indoor air of Households of urban area of Jammu.

MATERIALS AND METHODS

The study area was divided into eight sites: Site I (Households located near G.B. Pant Hospital, Nai Basti), Site II (Households in Commercial area, Jain Bazar), Site III (Households near National Highway I-A), Site IV (Households at Crossing, Satwari), Site V (Households in residential area but near Water Body, Jullaka Mohalla), Site VI (Households in residential area but near waste dumping site, Bhagwati Nagar), Site VII (Households in residential area but away from waste dumping site and big open drain, Sainik colony) and Site VIII (Households in residential area but near big open drain, Bakshi Nagar). At each site two households were selected randomly and in each household sampling of SPM and bacteria (CFU/ m^3) was done twice (i.e. once during July–Sept.2008 and once during Oct.–Dec.2008) at three sub sites i.e. Bedroom, Kitchen and Drawing room. Average value of each parameter with standard deviation for an average household at each site was compiled from data of twelve readings in a period of six months. Correlation coefficients (r) between SPM and gram +ve coccus, SPM and gram+ve bacillus, SPM and total numbers of Gram+ve bacteria, SPM and gram +ve coccus, SPM and gram+ve bacillus, SPM and total numbers of Gram –ve bacteria, SPM and total number of Coccus, SPM and total number of Bacillus

at all sites of study area were calculated using Pearson product-moment method.

Air Sampling for SPM was done by using Handy Air Sampler Envirotech APM 821 for two hours at a height of 5 ft above the ground. SPM was determined by formula:-

$$\text{SPM } (\mu\text{g}/\text{m}^3) = \frac{(W_2 - W_1) \times 10^3}{T \times \frac{R_1 + R_2}{2}}$$

Where W_1 and W_2 = initial and final weight of filter paper, R_1 and R_2 = initial and final flow rate in cubic metre.

T = sampling time in minutes

Air sampling for bacteria was done using Handy Air Sampler Envirotech APM 821 for 10 min at a height of 5 ft above the ground using sterile impingers containing 6 ml of distilled water. Three Petri plates i.e. one with Nutrient agar (peptic digest of animal tissue, beef extract, yeast extract, sodium chloride, agar, pH 7.4±0.2), second with MacConkey agar (peptic digest of animal tissue, lactose, sodium taurocholate, neutral red, agar, pH 7.4±0.2) and third with BTB Lactose agar (proteose peptone, beef extract, lactose, bromo thymol blue, agar, pH 8.6±0.2) were inoculated with 2ml. of impinged water from each impinger in Laminar flow and incubated at 37°C for 48 hrs in bacteriology incubator.

The quantification of bacterial count was done by using the formula:-

No. of microbes per volume (l) =

$\frac{\text{No. of microbes collected by impinger of air (CFU}/\text{m}^3)}{\text{Volume of air}}$

Volume of air

No. of microbes collected by impinger = Sum total number of colonies in all the three plates

Volume of air = Sampling time x flow rate of air in litre per minute

Sampling of bacteria was also done directly by exposing Petri plates with above said media to ensure that all the existing bacteria have been impinged. Bacterial study from each colony was carried out using Gram's staining technique under immersion oil. A control set for each culture media was prepared and the colonies found growing on the culture Medias were subtracted from the respective exposed culture Medias.

RESULTS AND DISCUSSION

The analysis of data revealed that Households near Hospital (Site I) exhibited minimum indoor SPM of 659.09±253.23 $\mu\text{g}/\text{m}^3$ whereas Households near Water body (Site V) exhibited maximum indoor SPM of 1006.12±224.95 $\mu\text{g}/\text{m}^3$. The Drawing room of Households in residential area away from waste dumping site and big open drain (Site VII) exhibited minimum indoor SPM of 224.46 $\mu\text{g}/\text{m}^3$ whereas Kitchen of the household at Site II i.e. in Commercial area exhibited maximum indoor SPM of 1560.54 $\mu\text{g}/\text{m}^3$ (Table I).

The Household located in Site II i.e. in Commercial area

exhibited minimum 1216.66 CFU/ m^3 count of Gram +ve cocci whereas Households in residential area but near Water Body i.e. Site V exhibited maximum value of 2448.31 CFU/ m^3 count. Gram +ve bacilli exhibited maximum value of 2842.82± 2062.08 CFU/ m^3 in households located in Commercial area and minimum value of 1168.58±343.37 CFU/ m^3 in Households located near National Highway I -A (Table I).

The Households at site I i.e. Near Hospital exhibited minimum Gram-ve cocci count of 1172.05±327.32 CFU/ m^3 and Households located Near Water Body (Site V) exhibited maximum value of 2409.77±1474.20 CFU/ m^3 . Gram-ve bacilli exhibited maximum and minimum values of 2475.01±1006.48 CFU/ m^3 and 1137.40±477.47 CFU/ m^3 at Site V i.e. Households Near Water Body and Site I i.e. Households Near Hospital (Table I), respectively.

The total average count of bacteria in the indoor air exhibited minimum value of 5251.00±1164.81 CFU/ m^3 with 56.01% (2941.55±1021.10 CFU/ m^3) of Gram +ve and 43.98% (2309.45±795.27) of Gram -ve bacteria at Site I i.e. Households Near Hospital and maximum value of 9308.24±2789.84 CFU/ m^3 with 47.52% (4423.46±1161.31) of Gram +ve and 52.48% (4884.77±1952.09) Gram -ve bacteria at Site V i.e. Households Near Water Body. Overall analysis at different sites of study area revealed that Households in the study area exhibited average indoor SPM of 801.25±300.73 with range of 224.46-1560.54. Analysis of data further revealed that Gram +ve and Gram -ve bacteria exhibited average bacterial count of 3552.30±1653.86 with a range of 821.21-10555.54 and 3246.14±1729.75 with a range of 1135.41-9062.5, respectively (Table I).

The critical analysis of the data revealed that Site V i.e. Households Near Water Body and Site VII i.e. Households in residential area away from waste dumping site and big open drain exhibited high %age of Gram-ve bacteria whereas the rest of the Sites viz. Site I i.e. (Households located near Hospital), Site II (Households in Commercial area), Site III (Households near National Highway I-A), Site IV (Households at Crossing), Site VI (Households in residential area but near waste dumping site), and Site VIII (Households in residential area but near big open drain) exhibited high %age of Gram +ve bacteria.

Overall analysis of data revealed that Households near Hospital surprisingly exhibited minimum indoor SPM and also the minimum value of bacterial count which might be due to maintenance of best sanitation conditions whereas Site V (Households near Water Body) exhibited maximum indoor SPM as well as maximum value of bacterial count due to dumping of silting material on banks of water body (Table I). Subba Rao *et al.* (1988) and Subramanyam *et al.* (1991) while studying microbial air quality of Madras city reported that increase in concentration of SO_2 and NO_x decreased microbial

Table 1. Indoor SPM and bacteria in households at different sites in Jammu city.

| Households | Indoor SPM ($\mu\text{g}/\text{m}^3$) | No. of bacteria (CFU/ m^3) in Indoor Air | | | | | | Grand Total |
|------------|--|--|--|---|--|---|--|--|
| | | Gram+ve | | | Gram-ve | | | |
| | | Coccus | Bacillus | Total | Coccus | Bacillus | Total | |
| Site I | 659.09 \pm 253.23 (293.16-1054.85) | 1771.26 \pm 77 0.28(842.1-3176.46) | 1170.30 \pm 3 59.01 (718.75-1789.47) | 2941.55 \pm 1 021.10 (1883.33-4941.16) | 1172.05 \pm 3 27.32 (789.47-1529.41) | 1137.40 \pm 477.47 (555.55-2368.41) | 2309.45 \pm 7 95.27(1345.02-4268.41) | 5251.00 \pm 1164.81 (3744.78-7464.88) |
| Site II | 900.04 \pm 326.93 (446.22-1560.54) | 1216.66 \pm 48 1.80(473.68-1852.94) | 2842.82 \pm 2 062.08(1138888.88) | 4059.48 \pm 2 269.26 (2036.18-10555.54) | 1947.44 \pm 1 013.13(777.77-4219.51) | 1745.06 \pm 8 39.45(777.77-3809.52) | 3692.50 \pm 1 525.26 (2444.43-8020.03) | 7751.98 \pm 3060.77 (5346.72-14836.81) |
| Site III | 707.79 \pm 239.07 (292.39-1054.85) | 1746.64 \pm 97 2.87(555.55-3684.20) | 1168.58 \pm 3 43.37(833.33-1894.73) | 2915.22 \pm 1 218.83 (1449.66-5578.93) | 1243.18 \pm 4 48.01(546.70-29164.705) | 1316.27 \pm 6 32.67(555.55-2631.57) | 2559.45 \pm 1 046.92 (1620.36-4531.57) | 5474.67 \pm 1637.57 (3126.73-8736.81) |
| Site IV | 859.02 \pm 160.63 (586.33-1054.85) | 1616.65 \pm 73 9.15(555.55-3078.94) | 1593.16 \pm 7 83.97 (894.11-3411.75) | 3209.81 \pm 1 113.39 (2326.3-4993.98) | 1541.46 \pm 7 19.94(777.77-3000) | 1212.03 \pm 4 86.48(555.55-2368.41) | 2753.49 \pm 1 043.83 (1345.02-4268.41) | 5963.30 \pm 1277.56 (4423.51-7528.66) |
| Site V | 1006.12 \pm 224.9 5(624.21-1393.24) | 2448.31 \pm 11 52.39(789.46-3999.99) | 1975.15 \pm 7 08.64(1052.63-3000) | 4423.46 \pm 1 161.31 (2158.72-5541.66) | 2409.77 \pm 1 474.20 (833.33-6562.5) | 2475.01 \pm 1 006.48 (1291.66-4875) | 4884.77 \pm 1 952.09 (2799-9062.5) | 9308.24 \pm 2789.84 (5161.73-14687.5) |
| Site VI | 778.56 \pm 343.78 (293.16-1251.95) | 2222.82 \pm 10 86.80(842.1-3999.99) | 1455.54 \pm 6 68.35(718.75-3000) | 3678.37 \pm 1 371.31 (1883.33-5062.5) | 1343.71 \pm 5 43.27(833.33-2250) | 1802.34 \pm 8 85.62(647.77-3250) | 3146.06 \pm 1 294.37 (1489.15-5277.76) | 6824.42 \pm 2202.47 (3744.78-10562.5) |
| Site VII | 799.29 \pm 303.00 (224.46-1393.24) | 1608.55 \pm 11 22.34(240.34-3875) | 1790.72 \pm 9 68.35(368.42-3000) | 3399.27 \pm 1 807.76 (821.21-5250) | 1944.23 \pm 1 735.77 (468.75-6562.5) | 1740.98 \pm 1 173.23(666.66-4875) | 3685.21 \pm 2 584.58 (1135.41-9062.5) | 7084.47 \pm 4127.15 (2441.54-14687.5) |
| Site VIII | 700.12 \pm 403.86 (293.16-1560.54) | 1371.26 \pm 68 4.09(473.68-3176.46) | 2419.96 \pm 2 220.61(718.75-8888.88) | 3791.22 \pm 2 426.14 (1883.33-10555.54) | 1578.03 \pm 9 75.82(842.10-4210.51) | 1670.43 \pm 8 96.39(647.77-3809.52) | 3248.46 \pm 1 767.17 (1489.15-8020.03) | 7039.68 \pm 3445.68 (3744.78-14836.81) |
| Average | 801.25 \pm 300.73 (224.46-1560.54) | 1750.27 \pm 95 6.02(661.76-8888.88) | 1802.03 \pm 1 301.35 (473.68-3437.5) | 3552.30 \pm 1 653.86 (821.21-10555.54) | 1647.48 \pm 1 058.00 (546.29-6562.5) | 1637.44 \pm 8 98.41 (555.55-4875) | 3246.14 \pm 1 729.75 (1135.41-9062.5) | 6837.22 \pm 2869.86 (2441.54-14836.81) |

Site I (Households located near G.B. Pant Hospital, Nai Basti), Site II (Households in Commercial area, Jain Bazar), Site III (Households near National Highway I-A), Site IV (Households at Crossing, Satwari), Site V (Households in residential area but near Water Body, Jullaka Mohalla), Site VI (Households in residential area but near waste dumping site Bhagwati Nagar), Site VII (Households in residential area but away from waste dumping site and big open drain, Sainik colony) and Site VIII (Households in residential area but near big open drain, Bakshi Nagar).

content of air whereas increase in SPM values also increased the microbial content.

A significant correlation (r) was found between SPM and gram +ve coccus (+0.13 to +0.66), SPM and gram+ve bacillus (+0.04 to +0.70), SPM and total numbers of Gram+ve bacteria (+0.18 to +0.78), SPM and gram –ve coccus (+0.06 to +0.63), SPM and gram–ve bacillus(+0.33

to +0.82), SPM and total numbers of Gram –ve bacteria(+0.21 to +0.76), SPM and total number of Coccus (+0.21 to +0.73), and SPM and total number of Bacillus (+0.33 to +0.81) at all sites of study area (Table II). Mancinelli and Shulls (1978), Subramanyam *et al.* (1999) also observed positive correlation between bacteria and SPM while studying airborne bacteria in urban

Table 2. Correlation coefficients (r) of bacteria and spm at different sites of Jammu city.

| SPM in households at different Sites | Number of Gram+ve | | Total Number of Gram+ve | Number of Gram-ve | | Total Number of Gram-ve | Total Number of coccus | Total Number of Bacillus |
|--------------------------------------|-------------------|----------|-------------------------|-------------------|----------|-------------------------|------------------------|--------------------------|
| | Coccus | Bacillus | | Coccus | Bacillus | | | |
| SPM at Site I | +0.42 | +0.04 | +0.35 | +0.37 | +0.47 | +0.43 | +0.58 | +0.42 |
| SPM at Site II | +0.33 | +0.27 | +0.31 | +0.49 | +0.54 | +0.62 | +0.50 | +0.41 |
| SPM at Site III | +0.54 | +0.40 | +0.54 | +0.34 | +0.36 | +0.36 | +0.62 | +0.48 |
| SPM at Site IV | +0.41 | +0.15 | +0.18 | +0.44 | +0.39 | +0.46 | +0.36 | +0.33 |
| SPM at Site V | +0.31 | +0.22 | +0.44 | +0.06 | +0.33 | +0.21 | +0.21 | +0.40 |
| SPM at Site VI | +0.56 | +0.46 | +0.67 | +0.42 | +0.82 | +0.74 | +0.73 | +0.81 |
| SPM at Site VII | +0.66 | +0.70 | +0.78 | +0.51 | +0.57 | +0.61 | +0.60 | +0.75 |
| SPM at Site VIII | +0.13 | +0.51 | +0.50 | +0.63 | +0.82 | +0.76 | +0.60 | +0.69 |

Pearson product-moment correlation coefficients are given for each bacterial i.e. total number of gram+ve (coccus and bacillus), total number of gram-ve (coccus and bacillus), total number of coccus and total number of bacillus and SPM pair.

Site I (Households located near G.B. Pant Hospital, Nai Basti), Site II (Households in Commercial area, Jain Bazar), Site III (Households near National Highway I-A), Site IV (Households at Crossing, Satwari), Site V (Households in residential area but near Water Body, Jullaka Mohalla), Site VI (Households in residential area but near waste dumping site Bhagwati Nagar), Site VII (Households in residential area but away from waste dumping site and big open drain, Sainik colony) and Site VIII (Households in residential area but near big open drain, Bakshi Nagar).

environment.

The present study also revealed that bacterial count in indoor air is affected more by indoor sources of pollutants than outdoor sources of pollutants. The bacteria enter air and disperse into air from various sources. The survival of these bacteria is a function of weather and the chemical composition of air. There was a statistically significant correlation (r) between the total number of bacteria and the concentrations of suspended particulate matter. It is reasonable to assume that bacteria in the air are protected from harsh environmental conditions, such as drying, by some of the suspended particulate matter (Mancinelli and Shulls, 1978).

The fluctuation in the number of Gram +ve Cocci and Bacilli as well as number of Gram -ve Cocci and Bacilli exhibited no set pattern of increase or decrease as it also depends on relative humidity, temperature and chemical pollutants. Mancinelli and Shulls (1978), also reported significant correlation between relative humidity and total

number of bacilli but observed no correlation between relative humidity and total number of cocci.

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