MICROBIOLOGICAL QUALITY ASSESSMENT OF RAW SALAD VEGETABLE SOLD IN MINNA METROPOLIS, NIGERIA

JD Bala*, FA Kuta, NU Adabara, AS Adedeji, UM Oyedum and G Murtala

Department of Microbiology, School of Life Sciences, Federal University of Technology, Minna, Nigeria

Abstract

Vegetables are edible part of plants. A total of twenty five raw salad vegetables were collected and the microbiological assessment was made using pour plate method. The analysis was carried out on carrots, cucumber, cabbage, lettuce and tomatoes. The results obtained from this study revealed that the total heterotrophic viable bacterial counts, coliform counts and fungal counts for all the salad vegetables ranged from $1.4 \times 10^6 \text{ - } 6.2 \times 10^6 \text{ cfu/g}$, $1.1 \times 10^6 \text{ - } 3.3 \times 10^6 \text{ cfu/g}$ and $2.1 \times 10^3 \text{ - } 4.5 \times 10^5 \text{ cfu/g}$ respectively. The data were subjected to One Way Analysis of Variance (ANOVA) test which showed that there was significant difference ($p \text{ < 0.05}$) in the microbial load of each of the raw salad vegetables samples. The microbial isolates identified were *E. coli*, *Staphylococcus aureus*, *Bacillus subtilis*, *Klebsiella* sp., *Pseudomonas* sp., *Aspergillus niger*, *Mucor* sp., *Penicillium* sp., *Aspergillus flavus* and *Fusarium* sp. *Staphylococcus aureus* and *Aspergillus niger* were predominant. This suggests that salad vegetables used in this study are of public health concern because they harbours microorganisms that could be hazardous to human health. Hence consumers should practice appropriate hygiene during the preparation of salad for consumption.

Key words: Assessment, microorganisms, quality, raw salad, vegetables

Introduction

Vegetables are considered as the leafy outgrowth of plants or plants shoot used as food. These include those plants or plant part used in making soup or served as an integral part of main meal (Yusuf et al. 2004). Vegetables can also be regarded as the edible component of plants, such components includes leaves, stalk, roots, tubers, bulbs, flowers and seed (ICMSF- International Commission on Microbiological Specification for Foods, 1974). Vegetables are important protective food and highly beneficial for the maintenance of health and prevention of diseases. They contain valuable food ingredients which are essential for the proper function of the body. Vegetable contain various medicinal and therapeutic agent and are valued mainly for their high vitamin and mineral content (Yusuf et al. 2004). Studies have evaluated the association of vegetables consumption with the reduction of risk of specific diseases (Hung et al. 2004). The occurrence of microorganisms in vegetables may be expected to reflect the sanitary quality of the processing steps and the microbiological condition of the raw product at the time of processing. Vegetables contamination in the field has been recognized as a source of human infection. Many of the viruses, bacteria and protozoan on vegetables which have caused food poisoning are derived from human feces (Ho et al. 1989, Rosenblum et al. 1990).

Foods, microorganisms and humans are in a long and interesting association that developed long before the beginning of recorded history. Foods are not only of nutritional value to those who consume them but often are ideal culture media for microbial growth (Willey et al. 2008). Vegetable which is defined as “any of the various plants especially herbaceous plants used wholly or partly for food” (Thomas 1995). Some of these

*Author for correspondence: bala.jeremiah@futminna.edu.ng/ jerrybrown316@yahoo.com
vegetables are cooked while others are eaten raw such as carrots, cabbage and lettuce. Some of these vegetables are grown with application of manure, fertilizer, and irrigation water which may serve as possible sources for contamination of the vegetables (Buck et al. 2003). Minimally processed ready-to-eat vegetables consist of raw vegetables that have been washed, peeled, sliced, chopped and shredded. Salads may be served with or without dressings depending on the consumer. For many types of raw or minimally processed foods, the microbial load is often composed of mixed species (Schuenzel and Harrison 2002). The washing process of vegetables reduces the microbial load approximately 10 folds thereby prolonging the shelf life (Gracia-Gimeno and Zurera-Cosano 1997). The survival and growth of pathogens on fresh product like vegetables is influenced by the organism, the product and the environmental conditions in the field and there after including storage conditions (Birbeck 2004). Environmental conditions can greatly influence microbial population due to the presence of free moisture on leaves from precipitation, dew or irrigation which may promote survival and growth of microbial pathogens (Brock and Lindow 1999). During vegetable handling, the process of washing and shredding can transfer microorganisms. Pathogenic microorganisms are able to infiltrate cracks, crevices and intracellular spaces of food products. However, pathogenic microorganism of human origin may also be present in minimally processed vegetables as the minimal technological processing may be unable to remove the original contamination resulting from air, soil, water, insects, animals, workers, harvesting and transportation equipment. Certain fungi such as Aspergillus, Fusarium, and Penicillium sp. as commonly occurring filamentous fungi grow in vegetable and their growth may result in production of toxins known as mycotoxins, which can cause a variety of ill effect in human from allergic responses to immunosuppression and cancer (Pitt et al. 1998).

Microbial growth on raw vegetables can result in the formation of biofilms by spoilage microorganisms which could provide protective environment for pathogens and reduce the effectiveness of sanitizers and other inhibitory agents. For example Listeria monocytogenes in a multi species biofilm with Staphylococcus aureus has been reported to be essentially unaffected by treatment with 500 ppm chlorine (Ketchum 2002). Biofilms have been observed on numerous leaf surfaces including leaves of lettuce and cabbage (Morrisa and Monier 1997). Fresh cut vegetables are considered a potential hazard since the occurrence of pathogens cannot be excluded and the product is consumed without heating. Thus, ready-to-eat salad vegetables could be a potential pathogen source if not hygienically processed before consumption. So far to the best of our knowledge, efforts have been geared towards studying the nutritional constituents of salad vegetables. As such there seem to be dearth of information on the microbiota been documented proving that a well-developed understanding of these is needed. Therefore, this study represents one of the few studies in this area. This will provide an insight to the microbiological characteristics of salad vegetables so as to lay a foundation of the microbiological aspects of salad vegetables in order to enhance better understanding of the microorganisms associated with salad vegetables particularly some pathogenic microbes that could cause health hazard and human diseases. Thus the study was designed particularly to determine the quality and microorganisms associated with salad vegetables in Minna, Niger state, Nigeria.

Materials and Methods

Collection of samples

A total of twenty five different raw salad vegetables were obtained from five different vendors. The samples included cabbage, tomatoes, cucumber, lettuce, and carrots (5 each) which were all obtained from different road side vendors within Minna, metropolis, Nigeria. All the samples were collected in sterile plastic bags and transported to the laboratory for processing immediately after collection.
Methodology and preparation of samples

The pour plate method was used as described by Jolt (2003) where 1ml of the appropriate serially diluted sample was transferred into labeled, well cleaned sterile Petri dish and molten agar medium (20 ml) poured. Poured plates were swirled gently to allow for proper distribution of colonies.

Bacteriological analysis

Total heterotrophic viable bacterial count

The method of Jolt (2003) was used where a 10 fold serial dilution of the samples was carried out. One gram of each of the salad vegetable was aseptically weighed and added into a sterile test tubes containing 9 ml of sterile, distilled water (deionized water). The test tubes were well shaken to mix together the content. Serial dilution was carried out using the mixed sterile distilled water as diluents. The 10-fold serial dilution was made by aseptically transferring 1 ml of the mixed distilled water in the test tubes into sterile test tubes containing 9 ml of sterile, distilled water. This gave ten times dilution. Subsequent dilutions were made from the aforementioned dilution. About 1 ml of the sample was pipette out from the $10^{-6}$ and $10^{-8}$ dilution tube into well labeled petri dishes. Then 20 ml of the molten nutrient agar was added into each plate and swirled gently to allow for proper mixing. The plates were incubated for 24 h at 37°C. Then the colonies develop on the plates were counted using a colony counter and expressed as colony forming unit per gram (cfu/g). The sample from each vendor was examined in triplicates and the average was recorded. The colonies differing in size, shape and colour were selected from the different plates on nutrient agar and sub-cultured repeatedly to obtain pure isolates. Macconkey agar was used to determine coliform counts. The pure isolates were maintained on agar slant for further characterization and identification.

Mycological analysis

The fungal count was carried out by pipetting 1 ml of the serially diluted salad vegetable onto Sabouraud Dextrose Agar (SDA) containing 0.01% chloramphenicol. An appropriate dilution of $10^3$ was used using pour plate method as described by Jolt (2003). The plates were incubated for 3 days at 25°C.

Characterization and identification of isolates

Bacterial isolates

The characterization and identification of the bacterial isolates were carried out based on cell morphology, Gram's reaction and biochemical tests according to methods described by Jolt (2003) and Oyeleke and Manga (2008). The isolates were identified by comparing with those of known taxa using the schemes of Cao (1996).

Fungal isolates

The mould isolates were characterized based on the colour of aerial and substrate hyphae, type of hyphae, shape and kind of asexual spores, presence of foot cell, sporangiophore, conidiophores, and the characteristics of spore head. A small portion of the mycelia growth was carefully picked with the aid of a sterile inoculating needle and placed in a drop of lactophenol cotton blue on a microscopic slide and covered with a cover slip. The slide was examined under the microscope, first with ($\times 10$) and then with ($\times 40$) objective lens to detect the spores and some special structures of the fungi. The isolates were identified by comparing their characteristics with those of known taxa using the schemes of Domsch and Gams (1970).
Results

Microbial counts

The results obtained for total viable bacterial counts, total coliform counts and total fungi counts for all the salad vegetables obtained in the present study ranged from $1.4 \times 10^6$ cfu/g - $6.2 \times 10^6$ cfu/g, $1.1 \times 10^6$ cfu/g - $3.3 \times 10^6$ cfu/g and $2.1 \times 10^3$ cfu/g - $4.5 \times 10^5$ cfu/g respectively (Table 1).

Frequency of occurrence of microorganisms isolated from salad vegetables

Table 2 revealed the frequency of occurrence of microbial isolates. *Staphylococcus aureus* and *Aspergillus niger* had the highest frequency of occurrence of 24.0 and 18.5 respectively.

Table 1. Total Microbial counts of the isolates.

<table>
<thead>
<tr>
<th>Samples</th>
<th>Total coliform</th>
<th>Total viable</th>
<th>Total fungi</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cucumber</td>
<td>$1.3 \times 10^{10\text{ab}}$ - $3.3 \times 10^{6\text{a}}$</td>
<td>$2.3 \times 10^{6\text{bc}}$ - $6.2 \times 10^{6\text{a}}$</td>
<td>$2.1 \times 10^{3\text{c}}$ - $6.3 \times 10^{4\text{c}}$</td>
</tr>
<tr>
<td>Carrot</td>
<td>$1.1 \times 10^{6\text{c}}$ - $2.2 \times 10^{6\text{a}}$</td>
<td>$2.8 \times 10^{6\text{c}}$ - $3.0 \times 10^{6\text{c}}$</td>
<td>$1.8 \times 10^{5\text{b}}$ - $4.5 \times 10^{5\text{a}}$</td>
</tr>
<tr>
<td>Cabbage</td>
<td>$1.2 \times 10^{6\text{bc}}$ - $3.2 \times 10^{6\text{a}}$</td>
<td>$1.4 \times 10^{6\text{bc}}$ - $2.1 \times 10^{6\text{c}}$</td>
<td>$1.9 \times 10^{4\text{c}}$ - $1.2 \times 10^{5\text{b}}$</td>
</tr>
<tr>
<td>Lettuce</td>
<td>$2.1 \times 10^{6\text{a}}$ - $2.2 \times 10^{6\text{a}}$</td>
<td>$1.4 \times 10^{6\text{d}}$ - $2.8 \times 10^{6\text{a}}$</td>
<td>$5.0 \times 10^{4\text{c}}$ - $1.2 \times 10^{5\text{b}}$</td>
</tr>
<tr>
<td>Tomato</td>
<td>$2.0 \times 10^{10\text{ab}}$ - $3.1 \times 10^{6\text{a}}$</td>
<td>$1.8 \times 10^{6\text{bc}}$ - $2.0 \times 10^{6\text{c}}$</td>
<td>$2.1 \times 10^{3\text{d}}$ - $1.2 \times 10^{4\text{c}}$</td>
</tr>
</tbody>
</table>

Values on the same column with different superscript (a, b, c, d) are significantly different from each other ($p < 0.05$) while those with the same superscript (a, b, c, d) are not significantly different from each other ($p > 0.05$).

Table 2. Frequency of occurrence of microorganisms isolated from salad vegetables.

<table>
<thead>
<tr>
<th>Organisms</th>
<th>No of occurrence</th>
<th>Frequency of occurrence (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Klebsiella</em> sp.</td>
<td>04</td>
<td>7.40</td>
</tr>
<tr>
<td><em>Bacillus subtilis</em></td>
<td>08</td>
<td>14.8</td>
</tr>
<tr>
<td><em>Staphylococcus aureus</em></td>
<td>13</td>
<td>24.0</td>
</tr>
<tr>
<td><em>Escherichia coli</em></td>
<td>05</td>
<td>9.30</td>
</tr>
<tr>
<td><em>Aspergillus niger</em></td>
<td>10</td>
<td>18.5</td>
</tr>
<tr>
<td><em>Mucor</em> sp.</td>
<td>04</td>
<td>7.40</td>
</tr>
<tr>
<td><em>Aspergillus flavus</em></td>
<td>03</td>
<td>5.60</td>
</tr>
<tr>
<td><em>Pseudomonas</em> sp.</td>
<td>02</td>
<td>3.70</td>
</tr>
<tr>
<td><em>Fusarium</em> sp.</td>
<td>02</td>
<td>3.70</td>
</tr>
<tr>
<td><em>Penicillium</em> sp.</td>
<td>02</td>
<td>3.70</td>
</tr>
<tr>
<td><strong>Total</strong></td>
<td><strong>54</strong></td>
<td><strong>100%</strong></td>
</tr>
</tbody>
</table>
Discussion

The results of the present study revealed that there was remarkable bacteria and fungi contamination of different salad vegetable samples. The samples were contaminated with varying degrees of pathogenic bacteria and fungi. The bacterial isolates identified in the present study include; *Escherichia coli*, *Staphylococcus aureus*, *Pseudomonas* sp, *Bacillus subtilis* and *Klebsiella* sp. The bacterial count in general, ranged from $1.1 \times 10^6$ cfu/g - $6.2 \times 10^6$ cfu/g and the fungal count ranged from $2.1 \times 10^3$ cfu/g - $4.5 \times 10^5$ cfu/g. This is in agreement with the report of Adebayo et al. (2012) who also reported similar microbial counts. The high count of bacteria detected in salad vegetables in the present study may be due to improper handling during harvesting and transportation. In the present study, all the vegetables examined harbored *Staphylococcus aureus*, while other microorganisms include *Bacillus subtilis*, *Klebsiella* sp., *E. coli* and *Pseudomonas* sp.

Most strains of *Staphylococcus aureus* are known to be pathogenic due to the heat stable enterotoxin they produce. The presence of *Staphylococcus aureus*, a pathogenic organism of public health concern and significance in these vegetables might have contaminated the vegetables from source as a result of handling by farmers or retailers. During processing, contamination could arise by use of dirty hands or clothing by food handlers and utensils used in slicing or keeping the vegetables. Improper handling and improper hygiene might lead to the contamination of food and this might eventually affects the health of the consumers (Omemu and Bankole 2005, Mgbakor et al. 2011).

The detection of *E. coli* in the study revealed poor hygienic standard in the handling of these raw salad vegetables or it could also be from contamination during harvest, it can also be present in water using in washing the salad vegetables. Presence of *E. coli* indicates recent contamination by faecal matter and possible presence of other enteric pathogens known to be causative agents of food borne gastroenteritis and bacterial diarrhea disease (Adebayo et al. 2012).

The fungal isolates obtained in the present study include, *Aspergillus niger*, *Aspergillus flavus*, *Fusarium* sp, *Mucor* sp and *Penicillium* sp. The presence of these fungi in raw salad vegetables have been reported in previous studies (Akintobi et al. 2011, Al-Hindi et al. 2011). Generally, fungi are considered toxigenic or pathogenic (Al-Hindi et al. 2011). During refrigeration, some moulds may produce mycotoxins on raw salad vegetables (Al-Hindi et al. 2011). Pathogenic fungi, on the other hand, could cause infections or allergies (Monso 2004, Al-Hindi et al. 2011). *Aspergillus* sp are known to produce several toxic metabolites, such as aflatoxin, maflorforms, naphthopyrones and they can produce ochratoxins (OTA), a mycotoxin which is a very important toxin worldwide because of the hazard it poses to human and animal health thus extra care should be taken during personnel handling of these vegetable, harvesting, cleaning, sorting, packaging, transport and storage (Pitt et al 1998, Petzinger and Weidenbach 2002, Al-Hindi et al. 2011). The result of the present study shows that the salad vegetables were contaminated by potentially pathogenic microbes. Therefore, salad vegetables could be a source of infection to the consumers if not properly washed before consumption (Al-Hindi et al. 2011).

Conclusion

The results of the present study revealed that the salad vegetables were all contaminated by a wide variety of potential pathogenic microorganisms. Proper hand wash with warm water and soap before handling the vegetable should be encourage. The vegetable should be thoroughly washed with clean water during the process of preparation. Food handlers should be advised on the need for good hygiene and the use of potable water for the washing of these vegetables to reduce the microbial load as low as possible.
References


