



Microbiological quality of minimally processed vegetables commercialized in Brasilia, DF, Brazil

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Abstract

This study evaluated the microbiological quality of minimally processed vegetables commercialized in the city of Brasilia, DF, Brazil. A total of 32 samples of different vegetables were purchased from 10 supermarkets. In most samples (78.1%) the populations of psychrotrophic bacteria had a high count ranging from 10^6 to 10^8 CFU/g. Thermotolerant coliforms were found in all samples, with populations higher than 10^2 MPN/g in 15 samples (46.9%). After molecular analyses, *E. coli* was identified in 16 samples (50.0%) and *Salmonella* spp. in 4 samples (12.5%). *S. aureus* was found in 14 samples (43.8%), with counts higher than 10^3 CFU/g in 4 samples (12.5%). The results obtained in this study showed that 16 samples (50%) were unfit for consumption according to Brazilian legislation. These results indicated the need of adoption of better hygienic practices in the production of minimally processed vegetables to improve quality and microbiological safety.

Keywords: fresh vegetables; microbiological analyses; bacterial pathogens; food contamination; food safety.

Practical Application: Microbiological quality assessment of minimally processed vegetables marketed in the city of Brasilia.

1 Introduction

Fruits and vegetables are essential components of a human healthy diet and there are considerable evidences of the nutritional benefits associated with the regular consumption of fresh fruits and vegetables (Goodburn & Wallace, 2013; Maffei et al., 2013). Consequently, nowadays, there is a growing interest in healthy diets, which increased consumption of vegetables and caused an expansion of the market for minimally processed vegetables. These products meet the needs of their consumers because of changes in the human lifestyle and its tendency to convenience and spending less time on preparing food (Gurler et al., 2015; Santos et al., 2012).

Minimally processed vegetables can be obtained from the fresh products through selection, washing, peeling, cutting or slicing, sanitization, rinsing, drying and packaging (Gurler et al., 2015; International Fresh-cut Produce Association, 2015; Tresseler et al., 2009). These products are defined as any fresh vegetable or combinations of them that has been physically altered from its original form, but remains in a fresh state, in order to preserve its nutritional and sensory properties (Cenci, 2011; International Fresh-cut Produce Association, 2015; Paula et al., 2009).

During processing, the mechanical damage caused to cells limits the shelf life of minimally processed vegetables. They are more susceptible to contamination because cutting and slicing damage the natural protective barriers of the intact vegetables and form exudates rich in nutrients that may support the growth of microorganisms. Because minimally processed vegetables are susceptible to spoilage, they must be kept refrigerated to

1-5 °C for maintenance of freshness and quality (O'Beirne et al., 2014; Oliveira et al., 2011). Processors of fresh-cut produce and supermarket retailers work hard to ensure the products are always kept cold. The consumers should also keep the produce refrigerated when they bring it home. When the abuse in the cold chain occurs, it accelerates microbial growth and spoilage of the products. The variation of temperature in 5 °C may lead to doubling of growth rate of any foodborne pathogens contaminating these products (Cenci, 2011; International Fresh-cut Produce Association, 2015; Paula et al., 2009).

Minimally processed vegetables can be contaminated with foodborne pathogens via being exposed to contamination sources during agricultural production, harvesting and handling, minimal processing and by end-users after pack opening. The major sources of contamination are soil, water (animal or human faeces) and handling during pre or postharvest stages. The major sources of postharvest contamination are containers used for transporting the produce, human handling, processing and storage (Cenci, 2011; International Fresh-cut Produce Association, 2015; O'Beirne et al., 2014). Minimal processing operations, including anti-microbial treatments, cannot be relied upon to eliminate pathogens (Gurler et al., 2015; O'Beirne et al., 2014).

Minimally processed vegetables have become increasingly recognized as potential safety problem, since they are usually eaten raw, without washing or other decontamination procedures. The incidence of foodborne outbreaks caused by contaminated fresh vegetables has increased in recent years (Herman et al., 2015;

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Little & Gillespie, 2008; Lynch et al., 2009). Several studies have isolated foodborne pathogens from raw vegetables, such as *Salmonella* spp. (Herman et al., 2015; Sant'Ana et al., 2011) and *Escherichia coli* (Herman et al., 2015; Söderström et al., 2008).

In Brazil, minimally processed vegetables are widely available, commercialized all year round and generally considered safe to eat by consumers. These vegetables can be stored for several days as minimally processed, since handled, packaged and refrigerated properly (Cenci, 2011; International Fresh-cut Produce Association, 2015; Sant'Ana et al., 2011). Therefore, the aim of this study was to evaluate the microbiological quality of minimally processed vegetables commercialized in supermarkets of Brasília, DF, Brazil.

2 Materials and methods

2.1 Samples and microbiological analyses

Individual packs of 100-200 g of minimally processed vegetables (32 samples) belonging to different producers and different brands were collected from ten different supermarkets in the city of Brasília, DF, Brazil, from December 2015 to July 2017. All the samples were obtained in the original package, within the shelf life of up to 6 days, as declared on labels. The samples were immediately transported to the laboratory in a cool box and microbiologically examined within 1 h of sampling. The samples were analyzed in triplicate and results were expressed as mean. The list of minimally processed vegetables types are shown in Table 1.

All the samples were analyzed for each of the following microorganisms or microbial groups: total mesophilic and psychrotrophic bacteria, total and thermotolerant coliforms, *Staphylococcus aureus* and *Salmonella* spp. An amount of 25 g of each sample was diluted in 225 mL of 0.1% peptone water. Samples were homogenized and an initial 10^{-1} dilution was obtained. Then, serial dilutions of the homogenates were prepared in 0.1% peptone water (up to 10^{-5}).

For total mesophilic and psychrotrophic bacteria counts, serial dilutions of the samples were surface plated in Plate Count Agar (PCA) (HiMedia, USA), following incubation at 37 °C for 24 h for mesophilic bacteria and 8-10 °C for 7 days for psychrotrophic bacteria. The results were expressed by colony forming unit per gram (CFU g⁻¹). For detection of total and thermotolerant

coliforms, one mL of each dilution was transferred to three-tube series containing Lauryl Sulfate Tryptose (LST) (HiMedia, USA) with Durham tubes in its interior. Total coliforms were enumerated in Brilliant Green Bile Broth 2% (HiMedia, USA), incubated at 37 °C for 24 h and thermotolerant coliforms were determined in *E. coli* broth (EC) (Acumedia, USA) incubated at 45 °C for 24 h. The results were expressed by most probable number per gram (MPN g⁻¹).

For total *Staphylococcus aureus* counts, serial dilutions of the samples were surface plated in Mannitol Salt Agar (HiMedia, USA), following incubation at 37 °C for 48 h. The colonies were counted and sub-cultured in Mannitol Salt Agar tubes. The characteristic colonies of *Staphylococcus aureus* (yellow colonies with yellow zones, mannitol-fermenting) stained by Gram's Method to confirm Gram-positive cocci. The colonies of *S. aureus* were further confirmed through molecular analyses.

The detection of *Salmonella* spp. was carried out with pre-enrichment in Lactose broth and incubation at 37 °C for 24 h. Subsequently, 1 mL of this medium was inoculated in Selenite Cystine and Tetrathionate broths (HiMedia, USA) that were incubated at 37 °C for 24 h. Further, *Salmonella* Shigella Agar (SS) and Xylose-Lysine Deoxycholate Agar (XLD agar) (HiMedia, USA) were streaked with sterile loops carrying inoculums took from Selenite cystine and Tetrathionate broths. The plates were incubated at 37 °C for 24 h in order to isolate characteristic colonies of *Salmonella* spp. Triple sugar iron (TSI) (HiMedia, USA) was used for presumptive confirmation of colonies that were further confirmed through molecular analyses.

2.2 Molecular analyses

The potential pathogenic bacteria *S. aureus*, *E. coli* and *Salmonella* spp. were identified using the technique of polymerase chain reaction (PCR). For identification of *S. aureus* the primers SEC forward and reverse (Table 2) (Life Technologies, Brazil) specific for the staphylococcal enterotoxin C gene (SEC gene) were used. For identification of *Salmonella* spp. the primers InvA forward and reverse (Life Technologies, Brazil) (Table 1) specific for the invasion A gene (invA gene) were used. And for identification of *E. coli* the primers MalB forward and reverse (Life Technologies, Brazil) (Table 2) specific for the formation of acetaldehyde and ammonia from ethanolamine (MalB gene) were used.

For DNA extraction, the NucleoSpin Food[®] kit (Macherey-Nagel, Düren, Germany), was used, following the manufacturer's instructions. Extracted DNA was stored at -20 °C. PCR was performed in a reaction mixture of 25 µl final volume containing 2.5 µl of PCR buffer; 0.7 µl of MgCl₂; 1.5 µL of dNTP (2,5 mM); 0.5 µl of Taq DNA polymerase; 1.5 µL of each primer forward and reverse and 18.3 µl of Milli-Q water.

PCR amplification was performed with an initial denaturing step at 95 °C for 1 min, followed by a 35-cycle reaction (95 °C for 1 min and 60 °C for 1 min). A final extension step was undertaken at 72 °C for 1 min. All thermal cycling reactions were performed with Techne TC-512 thermal cycler (Bibby Scientific Inc., USA). Both negative and reagent controls were included in each PCR run. The reagent control consisted of

Table 1. Composition and number of minimally processed vegetables samples analyzed in this study.

Composition	Number of samples (%)
Sliced collard green (<i>Brassica oleracea</i> L.)	13 (40.6)
Salads ¹⁾	7 (21.9)
Sliced cabbage (<i>Brassica oleracea</i> L. var. <i>Capitata</i>)	4 (12.5)
Lettuce (<i>Lactuca sativa</i> L.)	3 (9.4)
Mix for yakisoba ²⁾	3 (9.4)
Alfalfa sprouts (<i>Medicago sativa</i> L.)	2 (6.3)

¹⁾ Composed by two or more of the following vegetables: lettuce (different varieties), watercress, chard, cabbage, carrot, arugula and tomatoes; ²⁾ Composed by cauliflower, carrot, broccoli, cabbage and chard.

all PCR components except for the template DNA. All of the amplified DNA were separated by electrophoresis at 100 V for 50 min in 1.5% (w/v) agarose gel and stained with ethidium bromide. Gels were visualized under UV light. A 100 bp DNA ladder was used as a molecular weight marker.

3 Results and discussion

Table 3 shows the counts of mesophilic and psychrotrophic bacteria (CFU g⁻¹), enumeration of total and thermotolerant coliforms (MPN g⁻¹) and counts of *S. aureus* (CFU g⁻¹) in the samples of minimally processed vegetables. The populations of mesophilic and psychrotrophic bacteria in the tested samples were above 10⁴ CFU g⁻¹ and most samples had a high count ranging from 10⁶ to 10⁸ CFU g⁻¹ (20 samples or 62.5% for mesophilic bacteria and 25 samples or 78.1% for psychrotrophic bacteria). Although microbial loads tend to be higher for fresh vegetables (Maffei et al., 2013), a reduction of the microbial load is expected after its sanitization (Internacional Comissão on Microbiological Specifications for Foods, 2002). Some of these bacteria can grow during storage, especially those that are psychrotrophic and when packaged products are not stored at the temperature (usually between 5 °C) recommended by the provider products (Cenci, 2011; International Fresh-cut Produce Association, 2015).

Oliveira et al. (2011) also reported high populations of psychrotrophic bacteria (average populations of 10⁷-10⁹ CFU g⁻¹) in 162 samples of minimally processed vegetables (100% of analyzed samples) acquired from supermarkets in the city of Ribeirão Preto, São Paulo, Brazil. Similarly, Santos et al. (2012) found psychrotrophic bacteria count >10⁶ CFU g⁻¹ in 108 salads (71.47% of analyzed samples) commercialized in Portugal.

Generally, mesophilic and psychrotrophic bacteria counts and enumeration of total coliforms are useful for indicating the shelf-life duration and microbial quality of foods (Internacional Comissão on Microbiological Specifications for Foods, 2002). According to Oliveira et al. (2011) high populations of psychrotrophic bacteria suggesting a short shelf life for the product and poor hygienic quality, likely due to the use of highly contaminated raw material, lack of good hygienic practices during processing and/or inadequate temperature of storage.

Most samples of minimally processed vegetables (22 samples or 68.8% of analyzed samples) had enumeration of total coliforms higher than 10³ MPN g⁻¹. Similarly, Oliveira et al. (2011) reported the majority of samples of minimally processed vegetables (132 samples or 81.5% of analyzed samples) sold in supermarkets in the city of Ribeirão Preto with populations above 10³ MPN g⁻¹ for total coliforms. Silva et al. (2007) also reported high counts of total coliforms (>10³ MPN g⁻¹) for the 28 samples of minimally processed vegetables sold in supermarkets in the city of Porto Alegre.

In this study, thermotolerant coliforms were found in all samples of minimally processed vegetables, with populations higher than 10² MPN g⁻¹ in 15 samples (46.9%). The maximum value allowed by the Brazilian legislation for thermotolerant coliforms is 10² MPN g⁻¹ for fresh vegetables (Brasil, 2001). Thermotolerant coliforms may efficiently indicate failures in sanitization process, because they are associated with fecal contamination and they can indicate potential contamination by enteric pathogens (Paula et al., 2009; Silva et al., 2007).

Oliveira et al. (2011) reported that 45.6% of the samples analyzed (74 samples out of a total of 162) of minimally processed vegetables sold in supermarkets in the city of Ribeirão Preto exceeded the acceptable limit for thermotolerant coliforms of

Table 2. Primers sequence and size of amplified products of PCR.

Primer	Sequence 5' - 3'	Amplified product	Bacteria
SEC-foward	TTTACACCCAACGTATTAGCAGA	401bp	<i>S. aureus</i>
SEC-reverse	TCCCATTATCAAAGTGGTTTCC		
InvA-foward	GCTGATGCCGGTGAATTAT	445 bp	<i>Salmonella</i> spp.
InvA-reverse	CGACAAGACCATCACCAATG		
MalB-foward	TCTATGGGCTGTGACTGCTG	113bp	<i>E. coli</i>
MalB-reverse	GGCATCCCCATGATGTAGTT		

Table 3. Prevalence of microbial counts in the samples of minimally processed vegetables.

Count interval ^a	Mesophilic Bacteria n (%)	Psychrotrophic bacteria n (%)	Total coliforms n (%)	Thermotolerant coliforms n (%)	<i>S. aureus</i> n (%)
ND	-	-	-	-	18 (56.3)
10 ¹ -10 ²	-	-	3 (9.38)	17 (53.1)	2 (6.25)
10 ² -10 ³	-	-	7 (21.9)	11 (34.4)	8 (25.0)
10 ³ -10 ⁴	-	-	22 (68.8)	4 (12.5)	3 (9.38)
10 ⁴ -10 ⁵	7 (21.9)	2 (6.25)	-	-	1 (3.12)
10 ⁵ -10 ⁶	5 (15.6)	5 (15.6)	-	-	-
10 ⁶ -10 ⁷	7 (21.9)	12 (37.5)	-	-	-
10 ⁷ -10 ⁸	13 (40.6)	13 (40.6)	-	-	-

Count interval^a = expressed in CFU g⁻¹ or MPN g⁻¹; n (%) = number and percentage of positive samples; ND = not detected.

10^2 MPN g^{-1} . Paula et al. (2009) reported that 87.5% of the samples analyzed (42 samples out of a total of 48) of minimally processed vegetables (spinach, green beans, pumpkin and lettuce) sold in supermarkets in the city of Brasília, DF, Brazil, had populations of thermotolerant coliforms above 10^2 MPN. g^{-1} .

After molecular analyses, *E. coli* was found in 16 samples (50.0%) (Table 4). Oliveira et al. (2011) detected *E. coli* in 53.1% of the samples of minimally processed vegetables commercialized in city of Ribeirão Preto and Silva et al. (2007) detected *E. coli* in 8 samples (28.6%) of minimally processed vegetables commercialized in city of Porto Alegre.

The polymerase chain reaction (PCR) was used to amplify DNA sequences from the MalB gene of *Escherichia coli*. The MalB gene is specific for the formation of acetaldehyde and ammonia from ethanolamine. Candrian et al. (1991) used the MalB gene to detection of *E. coli* strains obtained from samples of Swiss-Italian soft cheese made from unpasteurized milk. Wang et al. (1997) also used the gene MalB to detection of *E. coli* in seafood samples. All *E. coli* strains tested yielded the specific DNA fragment (MalB gene) and no amplification products were obtained with other *Enterobacteriaceae* tested (*Enterobacter* spp., *Salmonella* spp. and *Yersinia enterocolitica*).

In this study, *Salmonella* spp. was detected and confirmed by PCR in 4 samples (12.5%) (Table 4). According to Brazilian legislation (Brasil, 2001), the presence of *Salmonella* in fresh vegetables is unacceptable (absence/25 g) and may represent a risk for consumers. The contamination rate of the samples from this study was higher than that observed by Oliveira et al. (2011) who found only 2 samples (1.2%) of minimally processed vegetables contaminated with *Salmonella* spp. Nguz et al. (2005) reported similar results and found an incidence of 13.3% of samples (8 samples out of a total of 60) of fresh mixed cut vegetables produced in Zambia contaminated with *Salmonella* spp. Gurler et al. (2015) reported 8 samples of mediterranean salads (14%) and 6 samples of sezar salads (12%) contaminated with *Salmonella* spp. These samples were collected from different restaurants, cafes and supermarkets in Turkey.

The pathogenic *Salmonella* spp. strains were identified by PCR used to amplify DNA sequences from the InvA gene. *Salmonella* pathogenicity depends on a variety of virulence factors that help the pathogen in adhesion and invasion mechanisms. Invasion gene (invA) exists in the majority of *Salmonella* strains and is related to intestinal mucosa invasion (Chuanchien et al., 2010; Fluit, 2005).

Foodborne diseases caused by *Salmonella* spp. and *E. coli* are usually associated with the consumption of animal products or cross-contamination by animal products. Contamination of vegetables with bacterial pathogens like *E. coli* and *Salmonella* can happen by animal manure used for fertilizer, water used for irrigation and animals that traverse vegetable fields. Contamination can also occur during processing (Herman et al., 2015; Nguz et al., 2005). Therefore, the potential contamination of vegetables by *E. coli* and *Salmonella* should be considered as these bacterial pathogens can survive and be internalized into plant tissues, rendering many disinfection treatments ineffective (Herman et al., 2015; O'Beirne et al., 2014).

According to Herman et al. (2015) during 1973-2012 in USA, 606 foodborne diseases and outbreaks had a leafy vegetable implicated (162 outbreaks with a simple leafy vegetable as the vehicle and 444 outbreaks with a leafy vegetable-based salad as the vehicle). The bacteria pathogens that most often caused leafy vegetable-associated outbreaks were Shiga toxin-producing *Escherichia coli* (STEC) and *Salmonella*. The STEC serogroup that majority caused the outbreaks was O157 (45 outbreaks, 94% of confirmed STEC outbreaks).

In this study, *Staphylococcus aureus* was found in 14 samples (43.8%) of minimally processed vegetables, with counts higher than 10^3 CFU g^{-1} in 4 samples (12.5%) (Table 3). The maximum value allowed by the Brazilian legislation for *S. aureus* is 10^3 CFU g^{-1} for fresh vegetables (Brasil, 2001). After molecular analyses, *S. aureus* producing the enterotoxin C gene was identified in 13 samples (Table 4). Nguz et al. (2005) reported an incidence of 60% (12 samples out of a total of 20) of *S. aureus* in fresh-cut mixed vegetables produced in Zambia. Seo et al. (2010) found an overall incidence of 11.3% of *S. aureus* in the 355 analyzed samples of minimally processed vegetables in Korea. At least 13 of 129 mixed salads samples (10.1%) and 19 of 112 sprouts (broccoli, alfalfa, clover and soybean) samples (17.0%) were contaminated with *S. aureus*.

The growth of *S. aureus* in food produces several enterotoxins that cause nausea, vomiting and diarrhea in consumers. Staphylococcal food poisoning is dependent on levels of toxin produced by *S. aureus* during growth. Staphylococcal enterotoxins (SEs) are a group of heatstable, pepsin-resistant enterotoxins belonging to a large family of pyrogenic toxin super-antigens encoded by phage (SEA), chromosome (SEB and SEC) or plasmid genes (SED) (Argudín et al., 2010; Seo et al., 2010). Enterotoxin C (SEC) is divided into subtypes C1, C2 and C3 and is an important cause of foodborne intoxication (Tamarapu et al., 2001).

Table 4. Occurrence of *E. coli*, *S. aureus* and *Salmonella* spp. in samples of minimally processed vegetables after PCR.

Sample	Total of samples	<i>E. coli</i>	<i>Salmonella</i> spp.	<i>S. aureus</i>
Sliced collard green	13	7	3	5
Salad	7	5	-	5
Sliced cabbage	4	1	1	2
Lettuce	3	1	-	-
Mix for yakisoba	3	1	-	1
Alfalfa Sprouts	2	1	-	-
Total	32	16	4	13

Table 5. Samples of minimally processed vegetables unfit for consumption.

Microorganisms above the limit* and/or presence of <i>Salmonella</i>	Sliced collard green	Salad	Sliced cabbage	Mix for yakisoba
Thermotolerant coliforms	4	3	2	1
Thermotolerant coliforms and <i>Salmonella</i> spp.	2	-	-	-
<i>S. aureus</i> and <i>Salmonella</i> spp.	-	-	1	-
Thermotolerant coliforms, <i>S. aureus</i> and <i>Salmonella</i> spp.	1	-	-	-
Thermotolerant coliforms and <i>S. aureus</i>	2	-	-	-
Total	9	3	3	1

*according to Brazilian legislation: Thermotolerant coliforms $>10^2$ MPN g⁻¹, *S. aureus* $>10^3$ CFU g⁻¹ and presence of *Salmonella* spp.

It is important to note that in case of minimally processed vegetables, *S. aureus* is a pathogen known to be carried mainly by food handlers and therefore good hygiene practices must be implemented by producers and processors in order to prevent *S. aureus* contamination. *Staphylococcus aureus* is a bacterium ubiquitously distributed in the environment and can be found on the skin and in the noses of up to 25% of healthy people and animals. The major route of transmission of *S. aureus* to food is through direct contacts with food handlers carrying the bacterium. *S. aureus* does not compete well with indigenous microflora in raw vegetables. However, *S. aureus* can grow in processed foods contaminated during preparation and storage under temperature-abused conditions (Argudín et al., 2010; Seo et al., 2010; Tamarapu et al., 2001).

The results obtained in this study showed that 16 samples (50%) were unfit for consumption according to Brazilian legislation (Table 5). Some samples presented more than one group of potentially pathogenic microorganisms such as a sample of collard green with presence of *Salmonella*, excess of thermotolerant coliforms and *S. aureus*, representing greater risk to consumer health.

4 Conclusion

In conclusion, this study showed that minimally processed vegetables may contain pathogenic bacteria and therefore could represent a risk to the consumers. A total of 32 samples were analyzed and the results showed that 16 samples (50%) were unfit for consumption according to Brazilian legislation. Good hygiene practices must be implemented by producers in order to minimize the risks of transmission of foodborne pathogens. Supermarket retailers and consumers must also guarantee that refrigerated conditions are always maintained.

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