

PREVALENCE OF PATHOGENIC BACTERIA IN COMMON SALAD VEGETABLES OF DHAKA METROPOLIS

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Abstract

Microbial quality of common salad vegetables (*viz.* carrot, cucumber, tomato and lettuce) collected from Dhaka metropolis was analysed to detect the presence of bacterial pathogens. The occurrence of huge numbers of fecal coliforms (1.0×10^4 - 4.09×10^6 cfu/g), *Escherichia coli* (1.0×10^4 - 5.0×10^8 cfu/g), *Staphylococcus aureus* (2.0×10^5 - 5.95×10^7 cfu/g), and *Listeria* spp. (1.5×10^6 - 6.5×10^7 cfu/g) were detected in all the tested samples. Interestingly, occurrence of viable but non-culturable (VBNC) bacteria was also noticed.

Introduction

Vegetables serve a major part of our food supply. Raw vegetables harbor a number of pathogenic microorganisms, which may be dispersed over the plants or appear as microcolonies embedded in the plant tissues (Beuchat 2002). During harvesting and transportation, raw vegetables may be bruised resulting in the release of plant nutrients, and thereby, providing substrates for microorganisms present on the surface of the vegetables to grow. In addition, the processing of fresh salad vegetables may alter or increase the number and type of pathogens present on the surface of the product. With a view of such exposure to pathogens, vegetables have been associated with the outbreaks of food borne disease in many countries (Alice 1997). Food borne illnesses can be caused mainly by microorganisms and/or their toxins. Cultivation of vegetables may largely account for such pathogenic contamination. Manures used to promote the growth of crops and vegetables contain a large number of pathogenic microorganisms including *Salmonella*, *Escherichia coli* O157:H7, *Bacillus anthracis*, *Mycobacterium* spp., *Brucella* spp., *Listeria monocytogenes*, *Yersinia enterocolitica*, *Clostridium perfringens*, *Klebsiella* spp. and *M. paratuberculosis* (Alice 1997). Therefore, a great risk towards public health is posed by the organic fertilizers applied in the fields. The major bacterial diseases caused are the various enteric diseases, diarrhoea, anthrax, salmonellosis, listeriosis, Crohn's disease, thrombocytopenic purpura, neurological disorders, arthritis, etc. (Cray and Moon 1995, Snowdon *et al.* 1989, Starutch 1991). Pathogens associated with untreated manure are assumed to enter into the food chain through crop. Thus, vegetables grown in such assistance of untreated fertilizers may play a significant role in showering pathogens to the consumers.

Therefore, an attempt was taken to assess the bacteriological quality, particularly pathogenic bacteria of fresh salad vegetables collected from several retail shops in Dhaka city.

Materials and Methods

A total of four different samples of common salad vegetables collected from different local markets and super shops of Dhaka City were used for microbiological analysis (Table 1). All the samples were collected according to the method suggested by American Public Health Association

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(APHA 1998). A portion of 25 g of each vegetable sample was aseptically weighed and buffer peptone water was added to make the final volume 100 ml. Serial dilutions were made up to 10^{-5} for plating (Cappuccino and Sherman 1996).

For isolation of fecal coliforms such as *Escherichia coli*, *Klebsiella* spp., *Staphylococcus aureus*, and *Listeria* spp., MFC agar, MacConkey agar, Manitol Salt Agar (MSA) and *Listeria* isolation media were used. After incubation at 37°C except for fecal coliforms (44.5°C) for 24 h, observations were made for the characteristic colonies (Cappuccino and Sherman 1996). The presence of *E. coli* was further confirmed by the appearance of bluish-black colonies with the production of green metallic sheen on Eosin-Methylene Blue (EMB) agar medium. During isolation, enrichment was performed for *Salmonella*, *Shigella* (in selenite cystine broth) and for *Vibrio* spp. (in alkaline peptone water) in order to avoid the false negative results due to the possibility of these bacteria to be in stressed condition or in the viable but non-culturable (VBNC) state in the samples, thereby decreasing the chance of their cultivability in the isolation media (Oliver 2005, Colwell 2000). For *Salmonella* and *Shigella* spp., Xylose Lysine Deoxycholate (XLD) plates were used while for *Vibrio* spp., Thiosulfate Citrate Bile Salt Sucrose (TCBS) agar plates were used. After incubation at 37°C for 24 h, characteristic colonies were detected and enumerated. Attempt was also taken to isolate *Clostridium* spp. from the vegetable samples (Francis *et al.* 1999).

Major biochemical tests such as Triple Sugar Iron (TSI), Motility Indole Urease (MIU), Methyl-Red (MR), Voges-Proskauer (VP), Citrate Utilization, Catalase test and Oxidase tests were carried out following the standard methods (Alfrad 2007, Cappuccino and Sherman 1996). All the experiments were performed in triplicate and the results were reproducible.

Results and Discussion

Fecal coliforms, *E. coli*, *Staphylococcus aureus* and *Listeria* spp. were found to be the most frequently proliferating pathogens in the vegetable samples (Table 1). *Klebsiella* spp. were found only in tomato (3.0×10^5). The load of *Vibrio* spp., *Salmonella* spp. and *Shigella* spp. were found to be nil; however, upon enrichment, the number of these pathogenic bacteria was found significantly higher. *Vibrio* spp. was estimated upon enrichment within a range of 2.0×10^4 - 8.3×10^7 cfu/g in carrot, lettuce and tomato samples, while *Salmonella* and *Shigella* spp. were found within a range of 1.0×10^3 - 3.1×10^7 cfu/g and 3.0×10^4 - 4.8×10^8 cfu/g respectively, in all samples. It is assumed that these pathogens were viable but non-culturable (VBNC) in the samples. A number of previous reports showed that *Vibrio cholerae*, *V. parahaemolyticus*, *V. vulnificus*, *Salmonella* and *Shigella* spp. may enter the VBNC state from which they are able to regain virulent properties after passaging in animal host (Colwell 2000, Huq *et al.* 2000, Olivier 2000, 2005). Thus, the assumed proliferation of the VBNC cells of *Vibrio*, *Salmonella* and *Shigella* spp. in fresh salad vegetables, as stated earlier, might be of significant interest in perspective of public health measure. All the isolates except *Listeria* spp. were confirmed through the biochemical tests (Table 2). It appears from the Table 1 that the samples from Shantinagar Bazar, Malibagh Bazar and Street site contained higher levels of microbial load than that of super shops. This might be due to the maintenance of hygienic condition with a relatively low storage temperature of the super shops. Prevalence of Staphylococci (2×10^5 to 5.95×10^7 cfu/g) in all the salad samples is indicative of health risk upon consumption of these raw vegetables. Presence of indicator organisms, *E. coli*, *Salmonella* spp., *Vibrio* spp., *Listeria* spp. and *Shigella* spp. revealed the possibility of spreading enteric diseases to the consumers. Similar results were also reported by Khan *et al.* (2008) and Nipa *et al.* (2011) from the similar vegetables.

Table 1. Bacterial load (cfu/g) of fresh vegetables.

Types of vegetables	Source	Fecal coliform	<i>Escherichia coli</i>	<i>Staphylococcus aureus</i>	<i>Listeria</i> spp.
Cucumber	Agora Super Shop	2.00×10^4	5.00×10^4	2.00×10^5	2.13×10^6
	Swapno Super Shop	1.00×10^4	5.65×10^5	3.42×10^6	1.57×10^6
	Malibag Bazar	6.90×10^5	3.32×10^7	6.55×10^6	2.55×10^6
	Shantinagar Bazar	5.42×10^5	4.42×10^7	8.42×10^6	5.43×10^6
	Street (Van) Sample	7.20×10^5	5.02×10^6	4.34×10^6	1.54×10^7
Carrot	Agora Super Shop	0	1.00×10^4	1.57×10^6	2.06×10^6
	Swapno Super Shop	0	1.40×10^5	3.11×10^6	3.48×10^6
	Malibag Bazar	4.60×10^5	1.23×10^7	6.84×10^6	6.72×10^6
	Shantinagar Bazar	8.33×10^5	2.72×10^7	9.53×10^6	8.42×10^6
	Street (Van) Sample	3.73×10^6	3.48×10^7	1.32×10^7	5.42×10^7
Lettuce	Agora Super Shop	3.00×10^4	6.50×10^5	4.31×10^7	1.50×10^6
	Swapno Super Shop	1.50×10^5	1.90×10^6	4.59×10^7	2.02×10^6
	Malibag Bazar	1.51×10^5	5.00×10^7	4.80×10^6	3.75×10^6
	Shantinagar Bazar	1.00×10^5	3.08×10^6	9.18×10^6	4.51×10^7
	Street (Van) Sample	3.00×10^6	5.00×10^8	5.95×10^7	5.00×10^7
Tomato	Agora Super Shop	0	2.20×10^5	5.00×10^4	-
	Swapno Super Shop	2.00×10^4	1.75×10^6	5.50×10^6	-
	Malibag Bazar	7.09×10^5	2.40×10^7	5.00×10^5	2.00×10^5
	Shantinagar Bazar	1.32×10^6	5.03×10^7	3.00×10^4	5.50×10^6
	Street (Van) Sample	4.09×10^6	1.23×10^8	5.02×10^6	6.50×10^7

Table 2. Results of biochemical tests of the pathogenic isolates.

Slant	TSI			Motility	Indole production	Urease activity	MR	VP	Citrate utilization	Catalase test	Oxidase test	Identified organism
	Butt	Gas	H ₂ S									
A	A	+	-	+	+	-	+	-	+	-	-	<i>Escherichia coli</i>
A	A	-	-	-	-	+	-	-	-	+	-	<i>Klebsiella</i> spp.
A	K	+	+	+	-	ND	+	-	+	ND	ND	<i>Salmonella</i> spp.
A	A	+	-	+	+	ND	+	-	-	ND	ND	<i>Shigella</i> spp.
+	+	-	-	-	-	-	+	-	-	+	-	<i>Staphylococcus aureus</i>
A	A	-	-	+	+	ND	+	-	+	ND	+	<i>Vibrio</i> spp.

A = Acidic reaction, K = Alkaline reaction, MR = Methyl red, VP = Voges-Proskauer, ND = Not done, + = Positive, - = Negative.

An important aspect of this study was the isolation of *Listeria* spp. from the salad vegetables which has been reported to represent an imperative risk to public health globally (Liu 2006, Little *et al.* 2007, Jeyaletchumi *et al.* 2010). Thus, the present study adds new insight to the existing knowledge on the microbiology of the commonly consumed fresh salads in Bangladesh. Another interesting facet from this study lies on the existence of VBNC state of the pathogens in the salad vegetables. Extensive identification of such cells would further unveil the factual implications of such bacteria on food borne epidemic outbreaks.

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