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Microbiological analysis and antibacterial activity of pear samples

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Present study attempted to isolate and quantify the microorganisms from the pear samples collected within Dhaka metropolis. Out of the five samples studied, total viable bacterial count and the total coliform count were near about 10^5 cfu/g and 10^4 cfu/g, respectively. Among the specific pathogens, prevalence of Salmonella spp. and Vibrio spp. were found only in two samples. Whereas Klebsiella spp, Pseudomonas spp. and Staphylococcus aureus were detected in all the samples studied. Fungal growth was also observed in each sample within the range of $8.8 \times 10^4 - 9.5 \times 10^4$ cfu/g. The study of antibiogram revealed that, most of the pathogens were resistant to more than one antibiotic. Interestingly the pear samples showed no anti-bacterial activity against any of the laboratory isolates such as, Escherichia coli, Pseudomonas spp., Salmonella spp. and Bacillus spp.

Key words: Pear, Pathogens; Fungal growth; Antibiogram; Anti-bacterial activity

The horticultural products particularly fruits contain vitamins, minerals, simple sugars and fibers that covers the needs of everyday people (1) and if their remedial and nutritional values are known then they can be used for both prevention and cure of diseases (2). Scientific data indicate that several degenerative diseases such as cardiovascular problems and various cancers can be prevented upon consumption of fruits and vegetables (3). However, as a result of unsuitable management and storage environment, the chance of microbial spoilage and diseases of a product increases by pathogenic and/or deteriorative microorganisms (4). The nature of fruits and vegetables is influenced by post harvest handling, transportation, storage and marketing because they are highly delicate products. [6] Indeed, intake of fresh-cut fruits and unpasteurized juices has lead to a number of outbreaks and cases of illness in the last years (5).

According to the report of WHO (1996) (7), the key factor of high mortality rates in developing countries is the infectious diseases. Several food borne pathogens such as *Escherichia coli*, *Salmonella* spp., *Listeria monocytogenes*, *Aeromonas* spp., *Staphylococcus* spp., *Streptococcus* spp., *Vibrio* spp. and *Pseudomonas* are responsible for various disease, morbidity and mortality (8).

The presence of Mycotoxins is also a matter of safety concern connected with foods. These are secondary metabolites of filamentous fungi; the most responsible and unsafe are species of the *Penicillium* and *Aspergillus* genera (8). They fall into the category of

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opportunistic microflora, with high possibilities of colonization since the harvesting period of horticultural products, when the pH of the tissue increased, the skin layers became soften and soluble carbohydrates build up and defense barriers is weaken (9). Now a day's treatment of various diseases is becoming difficult because of the emergence of drug resistant pathogens that may be introduced into the host because of the consumption of contaminated food (10-14). In recent years, many antibiotics have lost their effectiveness due to development of resistant strains as a result of the expression of resistance genes or spontaneous mutations within the microbial populations (15-19). This situation has contributed to the development of alternative antimicrobial drugs which are highly effective and non toxic, derived entirely from natural sources (20). Eventually, natural products with the healing properties has gained the interests of scientist community (21). Because of the presence of many active phytochemicals such as polyhenols, anthocyanins, flavonoids, terpenoids, carotenoids, cumarins, saponin and vitamins, fruits are becoming potential alternatives in the treatment of many diseases instead of using the antimicrobial agents (22-24).

Based on these facts and considerations, the current research focuses to ascertain and enumerate the pear spoiling microorganisms with their drug resistant traits and to reveal anti-bacterial activity of the samples tested.

MATERIALS AND METHODS

Selection of study area & collection of fruit samples for investigation. Samples were collected from some popular super shops from five areas of Dhaka city such as Rampura, Khilgao, Shantinagar, Moghbazar and Mailbagh areas. Samples were collected in different time intervals and transported to the laboratory as soon as possible according to the method suggested by American Public Health Association (25). For the identification & enumeration of pathogenic bacteria & fungi, at first 10g of each sample was taken, then blended with 90ml normal saline

(pH 7.8) & diluted up to 10^{-4} & then dilutions were used for plating purposes according to the standard guideline (26).

Enumeration of total viable bacteria & fungi. The enumeration was performed by using 0.1ml of each sample onto nutrient agar (NA) & Sabouraud Dextrose Agar (SDA) for the determination of total viable bacterial count (TVBC) and total fungal load respectively by the spread plate technique. The plates were incubated at 37 °C for 24 hours and at 25 °C for 48 hours for TVBC & total fungal load respectively.

Estimation of total coliform.. From the dilution 10^{-2} & 10^{-3} , the 0.1ml sample was spread onto MacConkey agar for the detection of total coliform. The plates were incubated at 37 $^{\circ}$ C for 24 hours.

Isolation of Salmonella spp., Shigella & Vibrio spp. The 1ml sample was transferred into 9ml of seleniote cysteine broth (SCB) and alkaline peptone water (APW) for the enrichment of Salmonella, Shigella and Vibrio spp. respectively & incubated at 37 °C for 6 hours. After enrichment the samples were diluted up to 10⁻⁴ & then 0.1ml of samples from 10⁻² & 10⁻³ were spread onto Salmonella-Shigella (SS) agar & Thiosulfate Citrate Bile Salt Sucrose (TCBS) agar followed by incubation at 37 °C for 24 hours for the detection of the colony characteristics. For the identification, the biochemical trait of the isolates was tested following standard biochemical methods (10, 11, 26).

Isolation of Staphylococcus spp., Pseudomonas spp. Staphylococcus & Pseudomonus spp were isolated from the Mannitol Salt Agar (MSA) & Pseudomonas agar respectively by spreading 0.1ml of the diluted samples on these media & then incubated at 37 °C for 24 hours.

Biochemical tests for the confirmative identification. Finally, the standard biochemical tests were performed to confirm the identification of all the pathogenic isolates found in all 5 types of pear samples by the previously described methods (26, 27).

Antibiotic susceptibility test. The pathogenic isolates were examined for antibiotic susceptibility traits (either drug resisistant or sensitive) by disc diffusion assay on Muller-Hinton agar (MHA) (Difco, Detroit, MI) against commonly used antibiotics following the standard protocol. Antibiotics used in the study are, Ampicillin (10 μ g), Amoxicillin (10 μ g), Ciprofloxacin (5 μ g), Ceftriaxone (30 μ g), Nalidixic acid (30 μ g), Imipenem (30 μ g), Erythromycin (15 μ g), Chloramphenicol (10 μ g), Trimethoprime-sulfomethoxazole (25 μ g), Gentamicin (10 μ g) & Piperacilline (10 μ g) (28).

Determination of antibacterial activity of the fruit sample. The antibacterial activity of the fruit samples was performed by using agar well diffusion method. Briefly, fruit blends were used directly on the Mueller-Hinton agar (MHA) media. At first, the bacterial pathogens (*Pseudomonas* spp, *Bacillus* spp, *Salmonella* spp, *E. coli*) were introduced evenly over the MHA separately using cotton swab, followed by making hole in the MHA by cork borer. Each of the blends was then introduced separately in the specified hole with a positive control (antibiotic disc-Gentamicin (GEN) 10µg) & a negative control (normal saline). Presence of clear zone around the sample solution (if any) indicated the presence of antibacterial activity (29-31).

RESULTS AND DISCUSSIONS

Most of the fruits are eaten raw and may lead to the outbreaks of human diseases that may put the overall public health at a serious risk. Various types of health related problems are associated with the proliferation of etiological agents as well as their drug-resistance properties (32). Along these lines, the present study portrays the (1) pathogenic load different types of pears consumed in Bangladesh, (2) Antibiogram profile of the pathogens, and finally (3) the antibacterial activities of these pear samples.

Prevalence of pathogenic microorganisms. Several previous studies showed that Escherichia coli, Salmonella spp., Shigella spp., Staphylococcus aureus, Vibrio spp. and fungi were common in a varieties of fruits and vegetables samples (13, 33). The load of Vibrio spp., Salmonella spp. and Shigella spp. were nil before enrichment; however, after enrichment Vibrio spp. was estimated within a range of 2.0×10^4 - 8.3×10^7 cfu/g while Salmonella and Shigella spp. were found in between 1.0×10^3 - 3.1×10^7 cfu/g and 3.0×10^4 - 4.8×10^8 cfu/g respectively. Another study showed that the Vibrio spp were found in different street food samples (10) by the margins of 5.2×10⁴ - 6.1×10⁶ cfu/g. Surprisingly total viable bacterial load (TVBC) in all the samples observed in this study was almost same, near about 10⁵ cfu/g whereas coliform, Pseudomonas and fungal count, all were within the range of 10⁴ cfu/g. Vibrio spp. was observed only in two samples within 8.0×10^4 - 9.0×10^4 cfu/g. On the other hand, range of the growth of Salmonella and Shigella spp. was $6.0 \times 10^4 - 8.0 \times 10^4$ cfu/g (Table 1). The presence of these pathogenic organisms revealed the possibility of spreading enteric diseases to the consumers. Presence of fungus indicates the presence of various Mycotoxins in the fruit samples that may pose severe threat to the human health. A study conducted by Renata Maria et al., (2011, 34) showed variable fungal growth in between 3.8×10^4 - 5.9×10^4 cfu/g. Overall fungal growth of the pear samples in this study was 8.8×10⁴- 9.5×10^4 cfu/g which is slightly higher than the previous report (Table 1).

So, this study showed the presence of different types of

TABLE 1. Load of various microorganisms in the pear samples

Sample No.	Microbiological Load cfu/ml									
	TVBC (cfu/g)	TCC (cfu/g)	Staphylococcus spp (cfu/g)	Salmonella spp (cfu/g)	Pseudomonas spp (cfu/g)	Fungi (cfu/g)	Vibrio spp (cfu/g)			
Sample-1	2.3×10 ⁵	4.0×10 ⁴	Nil	8.0×10 ⁴	6.0×10 ⁴	1.5×10 ⁴	Nil			
Sample-2	2.1×10^{5}	8.0×10^{4}	9.8×10^{4}	Nil	6.2×10^4	2.8×10^{4}	8.0×10^{4}			
Sample-3	2.1×10^{5}	9.0×10^{4}	8.9×10^{4}	Nil	3.4×10^{4}	8.8×10^{4}	Nil			
Sample-4	2.4×10 ⁵	3.5×10 ⁴	2.6×10^{4}	Nil	8.0×10^{4}	9.5×10^{4}	Nil			
Sample-5	2.4×10 ⁵	5.6×10 ⁴	6.2×10^4	6.0×10^{4}	9.0×10^{4}	6.2×10 ⁴	9.0×10^{4}			

TVBC = Total Viable Bacterial Count

TCC = Total Coliform Count

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Isolates	Klebsiella spp		Salmonella spp		Pseudomonas spp.		Staphylococcus spp.		Vibrio spp	
Antibiotics	R	S	R	S	R	S	R	S	R	S
AMP (10μg)	25%	75%	30%	70%	100%	0%	100%	0%	50%	50%
CIP (5µg)	50%	50%	60%	40%	100%	0%	50%	50%	70%	30%
PIP (10μg)	25%	75%	50%	50%	0%	100%	100%	0%	0%	100%
CEF (30µg)	50%	50%	25%	75%	0%	100%	ND	ND	35%	65%
AMO (10μg)	25%	75%	25%	75%	100%	0%	100%	1%	45%	55%
IPM (30µg)	0%	100%	0%	100%	0%	100%	25%	75%	20%	80%
CHL (10µg)	50%	50%	70%	30%	ND	ND	ND	ND	60%	40%
TMP-SUL(25µg)	25%	75%	60%	40%	25%	75%	25%	75%	75%	25%
GEN (10µg)	0%	100%	10%	90%	50%	50%	50%	50%	10%	90%
NALI (30µg)	75%	25%	80%	20%	0%	ND	ND	ND	75%	25%
ERY (15 µg)	50%	50%	60%	40%	ND	ND	25%	75%	55%	45%

TABLE 2. Antibiotic Susceptibility pattern of the isolates

AMP= Ampicillin, AMO= Amoxicillin, CIP= Ciprofloxacin, CEF= Ceftriaxone, NALI= Nalidixic acid, IPM= Imipenem, ERY= Erythromycin, CHL= Chloramphenicol, TMP/SUL= Trimethoprime-sulfomethoxazole, GEN= Gentamicin, PIP= Piperaciline. Sensitive- S; Registrant- R; not done- ND;

microorganisms with a high load in the pear samples. There are so many ways by which pathogens may come in contact with the fruits such as the crop land, organic fertilizers, irrigating water, packaging materials, transport vehicles etc. Besides, unhygienic personnel handling and processing of the fruits and their storage in such a condition which favors microbial growth might also account for such spoilage of fruits. The contaminating pathogens are responsible for various types of enteric diseases as well as serious intoxications in human health.

Antibiotic susceptibility patterns of pathogens found in pear samples. Drug resistance is a serious problem in these days that is becoming more and more risky for the global public health. Our study of antibiogram revealed, although some of the isolates were susceptible towards some antibiotics, several other antibiotics were proved ineffective, indicating the risk of the emerging resistant isolates causing health hazards. Almost every isolates were resistant for one or more antibiotics.

In studies conducted by Noor et al. (11, 12) showed that Staphylococcus spp. were highly resistant against ampicillin (93%), piperaciline (75%), amoxicillin (92%) and vancomycin. This result is quite consistent with this study (Table: 2). Intestinally, Klebsiella spp. was found to be sensitive against most of the drugs. But completely a different scenario has been found in a previous case, showing higher resistance of Klebsiella spp. against ampicillin (74%), ciprofloxacin (86%), piperaciline (88%), amoxicillin (72%) and ceftriaxone (97%) (11, 12). High resistance of Vibrio spp. against ampicillin, chloramphenicol, nalidixic acid. trimethoprimsulphametho xazole, tetracycline, cefotaxime has been shown by Acharjee et al. (10); whereas in this study variable susceptibility pattern were obtained against the different antibiotics for *Salmonella* and *Vibrio* isolates (Table 2). *Pseudomonas* spp were resistant against amoxicillin, ciprofloxacin and ampicillin whereas sensitive against ceftriaxone, piperaciline and imipenem (Table 2).

Antibacterial activity of pear samples. Chemically synthesized medicines may have many adverse side effects on our body but natural antimicrobials from the fruits can be used safely in near future as anti infective pharmaceutical products without creating any undesirable effects in our body. Antimicrobial activity of different fruits such as guava, orange, dragon fruit, Malta, apple, has been tested around the world different times (35). Interestingly, in the present study, antibacterial activity was not found in any one of the pear samples against the laboratory isolates of *Escherichia coli, Pseudomonas* spp., *Salmonella* spp. and *Bacillus* spp. However, the findings suggest that the study might be further extended by increasing the number of sample size to confirm and determine the antibacterial activity of the pear samples.

CONCLUSION

In fine, it can be concluded that the presence of a wide array of microorganisms with a huge load in the different pear samples is a matter of great concern for the consumers. The pathogenic microorganisms present in the commonly consumed pear fruit exhibited resistance against the regular antibiotics which is of significance from the view point of public health. Every possible measure should be taken by the different regulatory bodies to ensure the safe consumption of fruits.

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