



ISOLATION AND CHARACTERIZATION OF *ESCHERICHIA COLI* IN READY-TO-EAT FOODS VENDED IN ISLAMIC UNIVERSITY, KUSHTIA

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Abstract

Context: *Escherichia coli* is shed in the feces of warm blooded animals and humans and thus potential for public health. Detection and characterization of *E. coli* in the ready-to-eat (RTE) foods concerns due to their presence indicates fecal contamination of the food.

Objective: To identify, characterize and RFLP pattern analysis of *E. coli* isolated from RTE foods vended in Islamic University campus, Kushtia.

Materials and Methods: Fifty samples from four types of consumed foods in six student halls of residence, some temporary restaurants of Islamic University, Kushtia were assessed for bacterial contamination by standard methods. Identification and characterization of *E. coli* isolates were performed using IMViC tests. Genomic DNA was used to perform RFLP pattern analysis.

Results: Thirty seven out of 50 (74%) examined samples of RTE foods had *E. coli* contamination. The highest number of *E. coli* was isolated from vegetable oriented RTE foods (90.90%) and fish, meat and cereals samples were also significantly *E. coli* positive. RFLP profiling of two *E. coli* isolates were observed.

Conclusion: The results of this study provide evidence that some RTE foods had unsatisfactory levels of contamination with *E. coli*. Thus street vended RTE food could be important potential vehicles for food-borne diseases. Molecular characterization may be exploited to identify food borne pathogen among different species.

Keywords: Ready-to-eat foods, *Escherichia coli*, RFLP pattern.

Introduction

Food safety is an important part of public health, linking health to agricultural and other food production sectors (Schlundt 2002). Besides human sufferings and fatalities, the highest number of food borne outbreaks in recent years has devastating economic impact on food producer and processors. In addition, previously unknown food borne pathogens are constantly emerging (Newell *et al.* 2010) which are responsible for increased incidence of food borne diseases. These observations refer to street-vended foods. Familiarity, taste, low-cost and convenience are some of the appealing factors that make street foods popular as food source (Hanashiro *et al.* 2005). The street food industry plays an important role in developing countries (Muinde and Kuria 2005). Because of socioeconomic changes in many countries, this sector has experienced significant growth during the past few decades. They feed millions of people daily with a wide variety of ready-to-eat (RTE) foods and beverages sold and sometimes prepared in the streets or public places, relatively cheap and easily accessible (Barro *et al.* 2002).

The role of street food vendors in the transmission of diarrhoeal pathogens has been well-reported (Bryan *et al.* 1988). In Bangladesh, the presence of any pathogenic organisms in RTE food is a serious public health concern, as they do not receive any further treatment before consumption. Among bacteria, *Clostridium botulinum*, *Escherichia coli*, *Salmonella* spp., *Listeria monocytogens*, *Yersinia enterocolitica*, *Staphylococcus aureus*, *Shigella* spp., *Bacillus cereus* and *Campylobacter jejuni* comprise major part of food borne

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pathogens (Kumar *et al.* 2002). There is few data concerning the incidence of food borne diseases related to street foods in Bangladesh.

Among all food borne pathogen *E. coli* received much more attention. *E. coli*, a common inhabitant of the intestinal tract of humans and animals (Tannock 1995, Sørum and Sunde 2001), can be easily disseminated in different ecosystems through water supplies and the food chain (Perreten 2005). In addition, this microorganism can be implicated in human infections, mainly in immune-compromised patients (Huang *et al.* 2006) or when gastrointestinal barriers are violated, even normal 'nonpathogenic' strains of *E. coli* can cause infection.

About 98% of microbes found in food commodities are non-pathogenic (Kumar *et al.* 2002). Therefore it is necessary to characterize the target pathogen in order to develop a suitable diagnostic test to detect them. The conventional method to detect food borne pathogen is culturing technique, which include homogenization of food sample and enrichment in selective and nonselective media by plating to isolate pure culture. Finally phenotypic and genotypic characterization takes 3-4 days to confirm the results. Immunological and biochemical detection are time consuming; require pure culture and less sensitive over molecular method. DNA based method can deploy within mix culture of bacteria. Variation of DNA polymorphism among different bacterial species may exploit to identify food borne pathogen.

Therefore, the objective of this study was to determine the occurrence of *E. coli* in different types of RTE food samples collected from Islamic University campus, Kushtia, and to observe RFLP pattern.

Materials and Methods

Sample collection: Approximately 300 g of each fifty samples of RTE food of 4 categories (22 samples of cereals, 10 samples of meat, 7 samples of fish and 11 samples of vegetables) were collected between April 2009 and July 2009 from 6 student halls of residence and some temporary restaurants of Islamic university campus, Kushtia, Bangladesh. All samples were transported in sterile container to the laboratory and were tested within 24 h of collection. Ten grams of each food sample was mixed with one eighth strength Ringers Solution (Oxoid, Basingstoke, UK). The sample was homogenized with a blender (electric) at 6000 rpm for 5-10 minutes (10^{-1} dilution), followed by serial dilutions up to 10^{-6} dilution.

Standard coliform test and isolation of E. coli: Coliform counts were determined using the most probable number (MPN) method (Cappuccino and Sherman 1999, Harley and Prescott 2002). Fermentation tubes with appropriate quantity (10 ml) of lauryl tryptose broth medium were distributed with different strengths. The tubes were inoculated with 10 ml, 1 ml with 10 ml, and 1 ml and 0.1 ml amount of sample and incubated at 37°C for 24 h. All tubes of the presumptive test producing gas after 24 h of incubation, was further tested for conformation. A loopful inoculum from each culture showing production of acid and gas was transferred to Brilliant Green Bile Broth (Oxoid) and incubated for 48 h at 37°C and 44.5°C. Gas production at 44.5°C confirms presence of fecal coliform. Streaking on the eosine methylene blue (EMB) agar plate was done for further confirmation that was performed according to Cappuccino and Sherman (1999). One or more plates containing EMB agar medium were streaked from presumptive positive test tubes in such a way that discrete colonies may appear. The plates were incubated at 37°C for 24 h. Typical nucleated colonies with or without metallic sheen indicates positive results. A maximum of five suspected *E. coli* colonies from each sample (based on colony size, morphology and metallic green sheen on their surfaces) were selected. Further biochemical tests were done for the identification of *Escherichia coli* according to Cappuccino and Sherman (1999) and Coyle *et al.* (1985). IMViC test was performed to distinguish between *E. coli* and *Enterobacter aerogenes* (Cappuccino and Sherman 1999). Two isolates identified as *E. coli* were further characterized based on DNA polymorphism by RFLP.

Genomic DNA extraction. Standard and improved phenol chloroform method was used to extract genomic DNA (Neumann *et al.* 1992) with a few changes. Different samples were pre-treated properly for the collection of organism cell. In regard to pure bacteria culture, 1.4 ml bacteria suspension was collected by centrifugation at 16.1 rcf for 10 min at 4°C; precipitation was mixed well with 0.4 ml TE (10 mmol⁻¹ Tris-HCl, 1 mmol⁻¹ EDTA, and pH 8.0). After adding 50 µl of 10% SDS and 50 µl proteinase K, pellete was incubated at 42°C for 20 min in water bath. All nucleic acid came out of solution soon after adding 0.15 ml of 5M NaCl and 0.25 ml of ice cold isopropanol. The supernatant was transferred to a new tube and 0.7 vol of phenol/chloroform (1: 1) was added. After gentle mixing, the mixture was centrifuged at 16.1 rcf for 10 min at 4°C. The upper phase was transferred to a new tube containing 2/3 vol of isopropanol. The mixture was cooled at -20°C for 30 min and then centrifuged at 16.1 rcf for 10 min at 4°C. The resulting pellet was dissolved with 100 µl TE buffer after washing twice with 70% ice cold ethanol. The quantity and quality of the purified DNA were determined by measuring at A260 and by calculating the ratio of A260/A280, respectively by spectrophotometer. Finally, agarose gel electrophoresis of extracted DNA was performed using 0.8% UltraPure™ agarose (Invitrogen).

Restriction analysis. To get total 20 µl of volume, 1 µl of genomic DNA was mixed with 15 µl of distilled water and 2 µl of enzyme-assay buffer and finally with 2 µl of restriction enzyme (GeNei™) *i.e.* BamHI (10 U-µl), EcoRI (20 U-µl), EcoRV (10 U-µl), HindIII (20 Uv) and SacI (10 U-µl) added in different tubes. After incubation for 1 h at 37°C, complete digestion was checked on 3% agarose gel with using 100 bp-1 kb reference ladders.

Results

E. coli was isolated from 37 out of 50 (74%) food samples analyzed in this study; no *E. coli* isolates were obtained from the remaining 13 samples (including 7 samples of elaborated products derived from cereals). *E. coli* was detected in 10 out of 11 (90.90%) vegetables oriented RTE foods. All other examined fish (5 out of 7), meat (7 out of 10) and cereals (15 out of 22) RTE samples were significantly 71.42%, 70% and 68.18% contaminated with *E. coli* respectively.

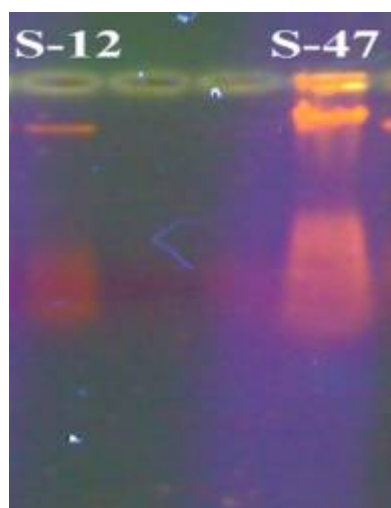


Fig. 1. Agarose gel electrophoresis pattern of the extracted DNA

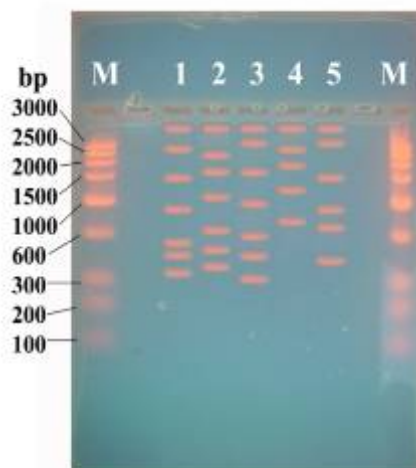


Fig. 2. RFLP pattern of one sample after electrophoresis. Lane1:BamHI, lane2:EcoRI, lane3: EcoRV, lane4:HINDIII, lane5: SacI

High yield genomic DNA extracted from two *E. coli* samples was obtained in present study. The A260 value of two DNA sample was 1.8199 and 1.7968. The A260/A280 ratio was 1.943 and 1.86 respectively. Some manipulations in the phenol/chloroform procedure were done. These were the addition of NaCl solution and ice cold isopropanol before phenol: chloroform extraction of samples. These two steps boosted the yield of high-quantity and high-quality genomic DNA and was amenable to further molecular characterization with RFLP. The integrity of the extracted DNA and restriction fragment pattern was checked by agarose gel electrophoresis (Figs. 1 and 2). The size of most digested DNA fragments ranged from 300 to 3000 bp (Fig. 2). RFLP digestion pattern might be the more powerful marker for molecular detection of pathogenic organism if coupled with PCR (e.g. RT-PCR). PCR amplification of the target gene with the appropriate oligonucleotide primer would be very specific for target organism. In future, the target gene of *E. coli* could be amplified and sequencing of the gene could reveal the better understanding of pathogenicity.

Discussion

A number of reports published on *E. coli* contamination of food. *E. coli* contamination (6.34%) was observed among 101 cooked and prepared food in a University centre in Argentina (Tessi *et al.* 2002). Another study conducted in Bangkok showed that coliform bacteria present in more than 50 percent of the food samples (Dawson *et al.* 1996). In the present study 74% food sample found as *E. coli* positive. The frequency of *E. coli* was found to be highest in vegetables. In Korea, *E. coli* was most frequently detected in convenient foods and 50% samples were found to be *E. coli* contaminated (Chung *et al.* 2010). Coliforms might be appeared in every phase of preparation; a case was reported (Seo *et al.* 2010) during investigation of the microbial quality of cooked soybean sprout salad. Presence of coliforms and *E. coli* in food may indicate fecal contamination which might be due to insufficient cooking, use of raw vegetables, cross contamination between raw and cooked food and contaminated ingredients (Eley 1992). So, presence of *E. coli* in 74% RTE food samples in the present study might be representing fecal contamination. People who depend on such food are often more interested in its convenience than in questions of its safety, quality and hygiene. Pathogenic bacteria including *S. aureus*, *E. coli* and *Salmonella* in restaurants would transfer to the cooked foods by contaminated staffs' hands or dishes (Nichols *et al.* 2002).

In our previous work, we reported that street foods in Dhaka city were extremely contaminated with microorganisms at an unacceptable limit especially the raw vendor items (Shaik *et al.* 2010). Total coliform was found to be present in all samples, indicating an alarming situation of health hazard. Bangladesh is an endemic zone for diarrhoeal diseases: every year, more than 5 % of death of children below 5 years of age is attributed to diarrhea (Arifeen *et al.* 2005). In Bangladesh, the predominant group of *E. coli*, associated with childhood diarrhea is ETEC, accounting for approximately 20% of all diarrhoeal cases (Qadri *et al.* 2005)

Conclusion

RTE foods vended in Islamic University campus, Kushtia had unsatisfactory levels of contamination with *E. coli*. Unhygienic practice may reveal the risk factors associated with contamination of post processing food. Food-borne disease is an urgent public health problem and requires rapid intervention. RFLP pattern analysis might be useful for molecular detection of pathogenic organism among different species if coupled with PCR. However, further detailed scientific study is necessary to develop rapid and easy detection method of pathogenic organism in food.

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