

Detection of Emerging Food Pathogens in Chicken Meat Using Multiplex Polymerase Chain Reaction

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Received: June 11, 2016

Accepted: July 14, 2016

Online Published: August 15, 2016

doi:10.5539/jas.v8n9p217

URL: <http://dx.doi.org/10.5539/jas.v8n9p217>

Abstract

A study was undertaken to develop a multiplex PCR (m-PCR) protocol for simultaneous detection of *Campylobacter jejuni* and *Listeria monocytogenes* in chicken meat. The extraction of DNA was carried out using commercial DNA extraction kit, Phenol Chloroform and boiling method. Samples with OD ratio (260:280) between 1.7 and 1.9 were considered good in terms of concentration and purity and were used for PCR amplification. DNA extraction kit and Phenol Chloroform method revealed good OD value were used for sample extraction process. PCR and m-PCR amplification was carried out using genus specific primers were designed by targeting its *Hyp* (500 bp) and *prfA* (290 bp) gene for *Campylobacter jejuni* and *Listeria monocytogenes* respectively. Electrophoresis of amplified PCR products and gel documentation revealed 500 bp and 290 bp in 2% Agarose. The multiplex PCR technique was standardized using the reference strains with the similar amplification procedure. The minimum detection level (sensitivity) by mPCR for *Campylobacter jejuni* and *Listeria monocytogenes* was found to be 0.2 ng/μl and 1.0 ng/μl of DNA in a reaction mixture (25 μl). The developed multiplex PCR technique could detect *Campylobacter jejuni* and *Listeria monocytogenes* upto 3×10^5 and 3×10^4 CFU/ml of artificially inoculated meat homogenate. Around 60 chicken meat samples were collected from different regions of chennai and were screened for the presence of *Campylobacter jejuni* and *Listeria monocytogenes*. All the samples screened were not positive either for *Campylobacter jejuni* and *Listeria monocytogenes*. The negative samples were further checked by culture methods and good correlation between these two methods was observed. Hence, the m-PCR technique developed in this study can be used as a rapid screening test for detection of *Campylobacter jejuni* and *Listeria monocytogenes* from chicken meat within 24 hours.

Keywords: chicken meat, *Campylobacter jejuni*, *Listeria monocytogenes*, multiplex polymerase chain reaction

1. Introduction

Recently many pathogens such as *Listeria monocytogenes*, *Campylobacter* spp., *Vibrio* spp., etc. have been identified as emerging meat borne diseases. *Campylobacter jejuni* is present in the gastro intestinal tract of all animals. Due to improper slaughtering method, the carcass may get contaminated with intestinal contents. Contaminated raw or undercooked poultry meats and/or by-products are particularly important to cause food-borne Campylobacteriosis in humans (CDC, 2005). The prevalence of contamination with *Campylobacter* species were approximately 60% in any portions of raw poultry meats and by-products except for fillets and hearts (Suzuki & Shigeki, 2008). *Listeria monocytogenes* is widely distributed and found in many food commodities. It is very persistent microorganism that survives on surfaces and equipment of food processing units in conditions of insufficient cleaning. Post processing contamination is the major source and cross contamination may also occur at the retail shop and also in products due to improper hygienic practices. Ingestion of uncooked meat contaminated during processing can produce infection. Contaminated raw meat was also a potential source for cross contamination to heat-treated or ready-to-eat products during processing or in the kitchen (Meyer, 2011). *L. monocytogenes* was found prevalently high in poultry (24.5%), intermediate in beef (24.4%) and less prevalent in pork (21.4%). Prevalence of *L. monocytogenes* in the eight isolated strains from ham and sandwiches was 37.5% and 25.0% respectively in Italy (Pesavento et al., 2010).

In earlier days, it was customary to sell the freshly slaughtered and dressed hot chicken meat with neither refrigeration nor storage facilities. Currently, increase in population (number of working women), changing life

style and food habits have propelled consumers to choose hygienic, packed, ready-to-cook meat and meat products. Hence, centralized meat processing units have been established to cater to this demand where there is possible risk of contamination of meat by many microorganisms. Therefore it is necessary to process and handle meat hygienically. Any discrepancy in maintaining the hygienic activities leads to increase in the microbial load in final product. Therefore all processors are opting for the implementation of HACCP in their plants to ensure food safety. It helps in identifying all biological hazards in various processing areas and monitors the critical points for which methods for rapid detection of pathogenic organism are required.

Therefore, there is a thrusting need to develop rapid and reliable method to detect the emerging food pathogens in meat while processing to deliver safe and quality meat and meat products to the consumers. Multiplex PCR is a type of PCR which enables simultaneous amplification of many target sites (template DNA) in one reaction by using more than one pair of primers for different organisms. It helps in minimizing the use of chemicals and also time needed for detecting single organism by normal PCR. Hence, the present study is undertaken for simultaneous detection of *Campylobacter jejuni* and *Listeria monocytogenes* from chicken meat by using multiplex polymerase chain reaction.

2. Materials and Methods

2.1 DNA Extraction

The reference strain of *Campylobacter jejuni* and *Listeria monocytogenes* (MTCC 657) was obtained from Department of Veterinary Public Health, Madras Veterinary College, Chennai and Institute of Microbial Technology (IMTECH), Chandigarh. The DNA was extracted by three methods –Bacterial DNA extraction kit method (Qiagen), Phenol-Chloroform-Isoamyl alcohol (25:24:1) extraction method and Boiling method. The genus specific primer was designed targeting *Hyp* gene: C-Forward: 5'-GGCGTTCATTTGGCGAATTTGAA-3', C-Reverse: 5'-CCGCTGTATTGCTCATAGGGA-3' and *prfA* gene: L-Forward: 5'-GAGCTATGTGCGATGCCA CTT-3' and L-Reverse: 5'-ATTAGCGAGCAGGCTACCGCAT-3', respectively.

2.2 Multiplex PCR (m-PCR)

A 25 µl of reaction mixture was set up in 0.2 ml PCR tube with following components such as master mix - 12.5 µl, forward primer (*Listeria*) - 1 µl, reverse primer(*Listeria*) - 1 µl, forward primer (*Campylobacter*) - 1 µl, reverse primer(*Campylobacter*) - 1 µl, template DNA - 2 µl and nuclease free water - 6.5 µl. The PCR amplification was carried out in Master Cycler Gradient Thermo cycler (M/s. Eppendorf, Germany) with the following cycling of initial denaturation at 94 °C for 5 minutes, followed by 30 cycles of denaturation (94 °C for 30 seconds), annealing (52 °C for 30 seconds) and extension (72 °C for 30 seconds) and subsequently a final extension at 72 °C for 7 minutes. The PCR product was subjected to electrophoresis in 2% agarose gel with ethidium bromide added at a concentration of 1 µg/ml of agarose. Electrophoresis was carried out in 1 X TAE buffer at 100 volts for 30 minutes. The gel was viewed under UV transilluminator and photographed in gel documentation system.

2.3 Minimum Detection Limit or Sensitivity of Multiplex PCR

DNA from reference strain was extracted and quantified using Biophotometer plus (M/s Eppendorf, Germany). The quantified DNA was serially 10 fold diluted in sterile nuclease free water and for each dilution PCR amplification was carried out. Sensitivity of m-PCR in terms of DNA concentration was determined. The highest dilution of the DNA showing a visible band in gel was taken as the detection limit for *Campylobacter jejuni* and *Listeria monocytogenes*.

2.4 Artificial Inoculation of *Campylobacter jejuni* and *Listeria monocytogenes* in Chicken Meat

Pellets obtained from overnight incubated Bolton broth and Brain Heart Infusion broth containing pure cultures of *Campylobacter jejuni* and *Listeria monocytogenes* (MTCC 657) respectively were separately suspended in sterile saline solution. The cell numbers were adjusted from 3×10^8 CFU/ml to 3×10^2 CFU/ml by doing serial 10 fold dilutions. The cell numbers in each dilution were confirmed by plate count method using Blood free *Campylobacter* Selective agar and *Listeria* Identification agar (PALCAM).

The chicken meat samples which were negative for *Campylobacter jejuni* and *Listeria monocytogenes* were confirmed by culture methods was used for artificial inoculation study. Both of these bacteria in respective tenfold dilutions from 3×10^8 CFU/ml to 3×10^2 CFU/ml (0.2 ml of each dilution) were inoculated together in 1.8 ml of meat homogenate obtained by homogenizing 25 gram of chicken meat in 225 ml of BPW. DNA was extracted before pre-enrichment of meat homogenate, by using Bacterial DNA Extraction Kit. The minimum detection level (sensitivity) for m-PCR in terms of number of organisms was determined. The highest dilutions

of the cultures showing visible bands in gel were taken as the detection limit for *Campylobacter jejuni* and *Listeria monocytogenes*.

2.5 Screening of Chicken Meat Samples

Around 60 chicken meat samples were collected from different regions of Chennai. These samples were screened for the presence of *C. jejuni* and *L. monocytogenes* by the m-PCR technique developed in this study. The meat homogenate obtained was then subjected to DNA extraction using Phenol-Chloroform-Isoamylalcohol mixture (25:24:1) and m-PCR analysis for the presence of *C. jejuni* and *L. monocytogenes*. The chicken meat samples, in which either *Campylobacter jejuni* and *Listeria monocytogenes* or both were detected by m-PCR, were further confirmed by culture methods.

3. Results and Discussion

3.1 DNA Extraction

Three methods of DNA extraction were tried, out of which the Phenol Chloroform Isoamyl alcohol (25:24:1) extraction method was found to be cost effective and the purity and concentration of DNA (Table 1) obtained by this method was almost equal to that of Real Genomics DNA extraction kit method. The purity of DNA (OD ratio, 260 nm:280 nm) was considered good when the values of OD ratio were between 1.7 and 1.9 (Stephenson, 2010). For boiling method, the purity of DNA was not considered good, the reason for not getting good purity may be due to presence of other compounds than DNA and the other materials are not properly separated or removed during extraction procedure.

Table 1. Mean \pm SE values of purity of DNA and concentration of DNA

DNA Extraction method	Purity (OD ratio)		Concentration(μ g/ml)	
	<i>C. jejuni</i>	<i>L. monocytogenes</i>	<i>C. jejuni</i>	<i>L. monocytogenes</i>
Qiagen Bacterial DNA extraction kit	1.85 \pm 0.01	1.86 \pm 0.02	876.67 \pm 0.92	653.83 \pm 0.87
Phenol Chloroform Isoamyl alcohol	1.78 \pm 0.02	1.85 \pm 0.02	602.50 \pm 0.99	501.83 \pm 0.79
Boiling method	1.09 \pm 0.01	1.06 \pm 0.01	950.17 \pm 0.48	800.83 \pm 0.60

Among the two extraction methods, Qiagen kit method was best because of higher concentration and purity of DNA which may be due to the presence of spin column which can easily remove or filter the other impurities leaving the DNA. Even though the kit method was considered as best, Phenol-Chloroform-Isoamyl alcohol (25:24:1) extraction method was cost effective. Hence Phenol-Chloroform-Isoamyl alcohol method was standardized and used for further screening of retail chicken meat.

3.2 Multiplex PCR (m-PCR) Standardization

Using the genus specific primers newly designed for *Campylobacter jejuni* and *Listeria monocytogenes* by targeting the *Hyp* gene and *prfA* gene, DNA were amplified which gave the product size of 500 bp and 290 bp respectively by PCR (Figure 1). The multiplex PCR technique was standardized to detect both *Campylobacter jejuni* and *Listeria monocytogenes* simultaneously in the same reaction tube (Figure 2). Electrophoresis performed in 2% agarose, consistently gave good results with 500 bp of *Campylobacter jejuni* and 290 bp of *Listeria monocytogenes*. The results of this study were congruent with that of Wang and Slavik (2005) who developed a multiplex PCR assay to detect *C. jejuni*, *E. coli* O157:H7, *S. typhimurium* and *L. monocytogenes* with amplified PCR products of sizes 159 bp, 252 bp, 360 bp and 450 bp respectively. Similar findings were also reported by Yamazaki-Matsune et al. (2007) who developed a multiplex PCR for different species of *Campylobacter*.

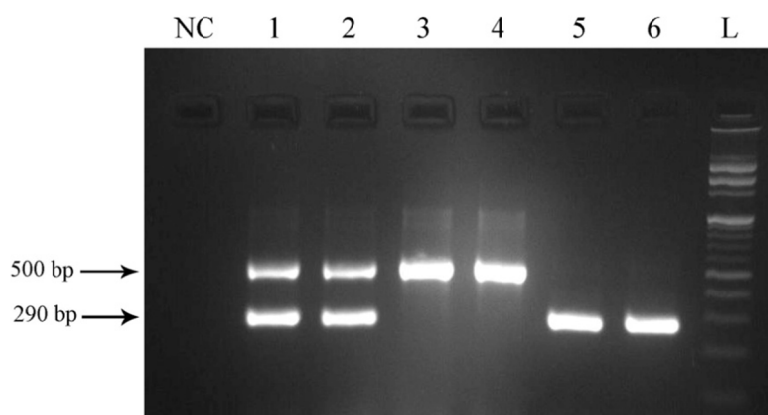


Figure 1. Standardization of PCR and multiplex PCR-Agarose gel electrophoresis of PCR products

Note. L: 100 bp DNA Ladder, 1-2: *Campylobacter jejuni* and *Listeria monocytogenes* revealing 500 bp and 290 bp mPCR product of *Hyp* and *prfA* gene, 3-4: *Campylobacter jejuni* revealing 500 bp PCR product of *Hyp* gene, 5-6: *Listeria monocytogenes* revealing 290 bp PCR product of *prfA* gene NC: Negative control.

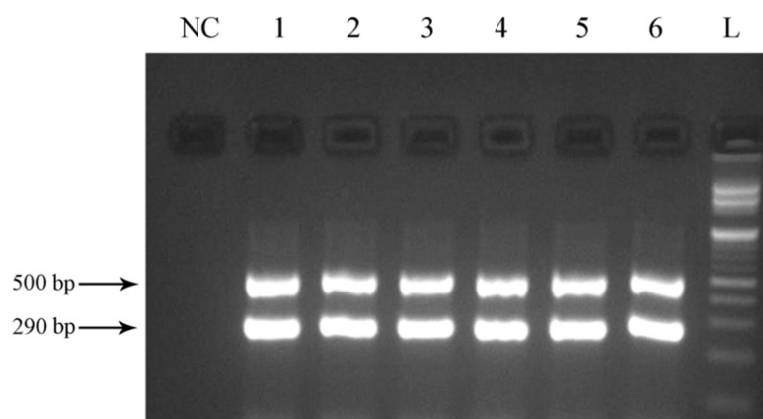


Figure 2. Standardization of m-PCR- Agarose gel electrophoresis of mPCR products

Note. L: 100 bp DNA Ladder, 1-6: *Campylobacter jejuni* and *Listeria monocytogenes* revealing 500 bp and 290 bp PCR product of *Hyp* and *prfA* gene, NC: Negative control.

3.3 Sensitivity of Multiplex PCR (m-PCR) for *Campylobacter jejuni* and *Listeria monocytogenes*

Diluted DNA of *Campylobacter jejuni* and *Listeria monocytogenes* were used to amplify in the same reaction tube and the sensitivity (minimum detection limit) of Multiplex PCR (m-PCR) for *Campylobacter jejuni* and *Listeria monocytogenes* was found to be 0.2 ng/μl and 1.0 ng/μl of DNA in a reaction mixture (25 μl) respectively (Figure 3). Related results were obtained by Keramas et al. (2003) who developed a multiplex PCR and detected DNA from *C. jejuni* and *C. coli* to a minimum level of 15-150 fg/μl. He also concluded that less sensitivity of bacterial detection was noticed in m-PCR when compared with PCR. The sensitivity in the study was lower when compared to that of Keramas et al. (2003).

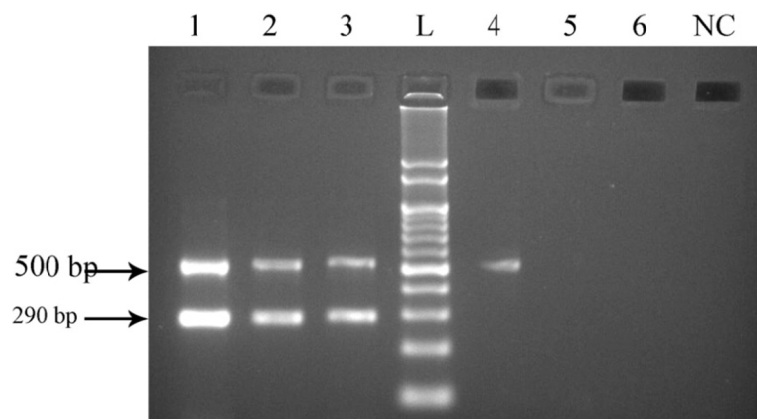


Figure 3. Agarose gel electrophoresis of m-PCR products for Minimum detection level of *Campylobacter jejuni* and *Listeria monocytogenes*

Note. L: 100 bp DNA Ladder, 1-6: PCR amplification performed with 200 ng/μl, 20 ng/μl, 2 ng/μl, 0.2 ng/μl, 0.02 ng/μl, 2 pg/μl of *Campylobacter jejuni* DNA and 100 ng/μl, 10 ng/μl, 1 ng/μl, 0.1 ng/μl, 0.01 ng/μl, 1 pg/μl of *Listeria monocytogenes* DNA respectively. Note no amplification with 0.02 ng/μl, 2 pg/μl of *Campylobacter jejuni* DNA and 0.1 ng/μl, 0.01 ng/μl, 1 pg/μl of *Listeria monocytogenes* DNA respectively. NC: Negative control.

3.4 Artificial Inoculation of *Campylobacter jejuni* and *Listeria monocytogenes* in Chicken Meat

Artificially inoculated reference strains of *Campylobacter jejuni* and *Listeria monocytogenes* (MTCC 657) from 3×10^8 CFU/ml to 3×10^3 CFU/ml, DNA was extracted from the meat homogenate and the developed m-PCR technique was used for detection purpose (Figure 4). The sensitivity of *Campylobacter jejuni* and *Listeria monocytogenes* were found to be 3×10^5 and 3×10^4 CFU/ml of meat homogenate respectively by the developed m-PCR technique. Gilbert et al. (2003) detected bacterial concentrations greater than or equal to 10^4 CFU/reaction for *C. jejuni*, *E. coli*, *Salmonella* and *L. monocytogenes*. Zhang et al. (2009) developed a m-PCR system for simultaneous detection of *Listeria monocytogenes*, *Staphylococcus aureus*, *Salmonella enterica* and *E. coli* O157:H7 with a sensitivity as low as 1 CFU/ml per 25 g food. Similarly Germani et al. (2009) reported that *Listeria* alone can be detected at 10^1 cells/ml in PCR, whereas with multiple pathogens in m-PCR they noticed a reduction in detection level of 10^6 cells/ml.

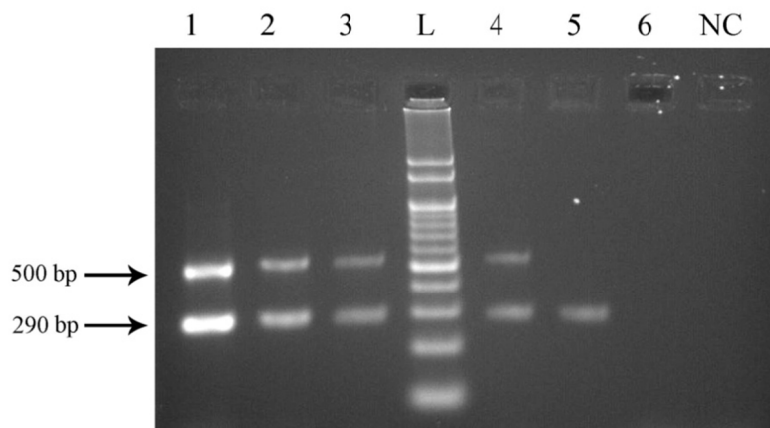


Figure 4. m-PCR amplification for chicken meat sample spiked with *Campylobacter jejuni* and *Listeria monocytogenes*

Note. L: 100 bp DNA Ladder; 1-6: PCR amplification performed with 3×10^8 CFU/ml, 3×10^7 CFU/ml, 3×10^6 CFU/ml, 3×10^5 CFU/ml, 3×10^4 CFU/ml, 3×10^3 CFU/ml of *Campylobacter jejuni* and *Listeria monocytogenes* respectively. Note no amplification with 3×10^4 CFU/ml, 3×10^3 CFU/ml of *Campylobacter jejuni* and 3×10^3 CFU/ml of *Listeria monocytogenes*; NC: Negative control.

3.5 Sequencing

PCR products were confirmed by sequencing the samples from Ocimumbio Solutions (P) Ltd, Hyderabad. Then the data were analysed by using Chromas Pro sequencing analysis software and the chromatograph for separate organism with forward and reverse primer (Figures 5, 6, 7 and 8).

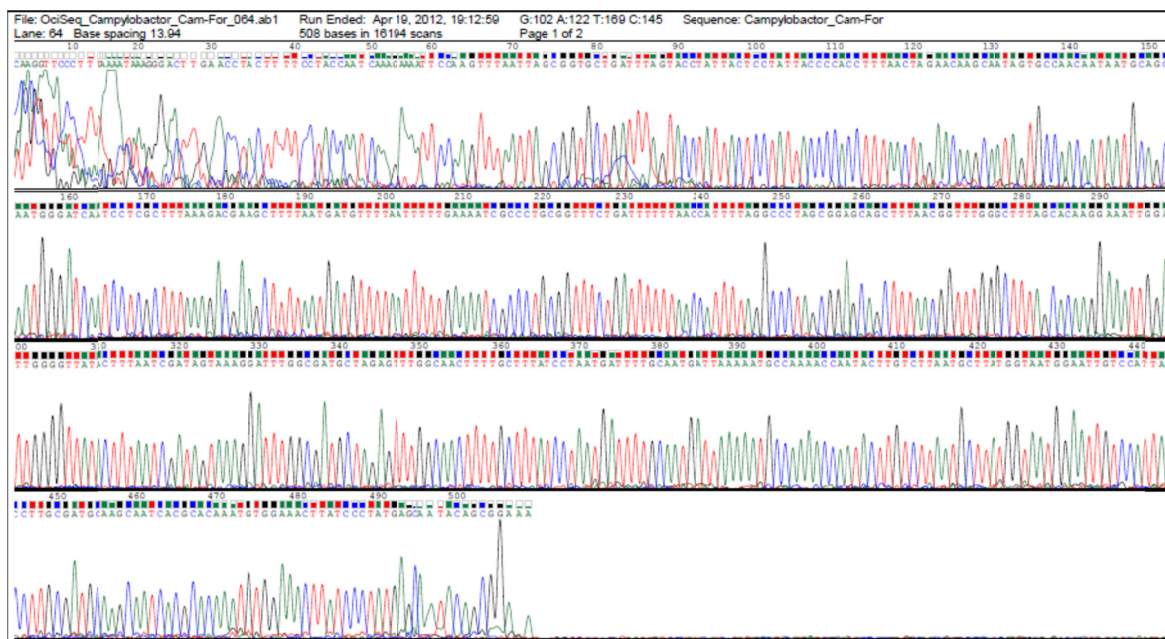
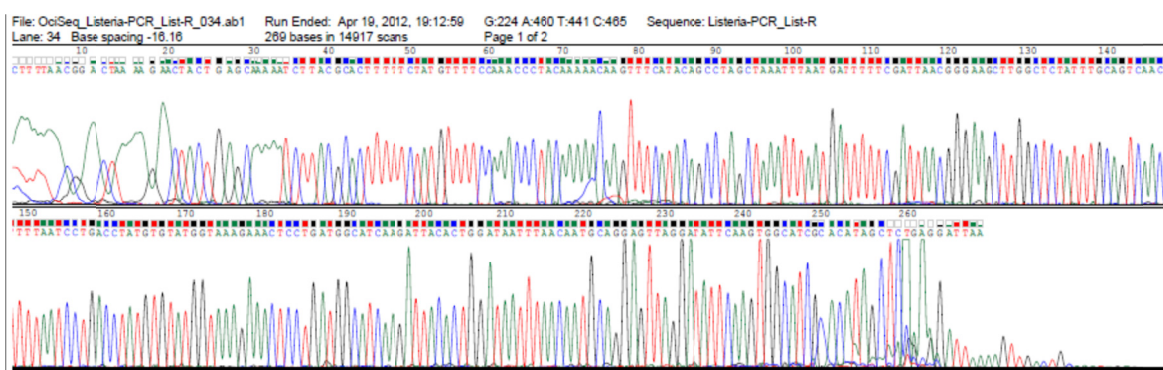
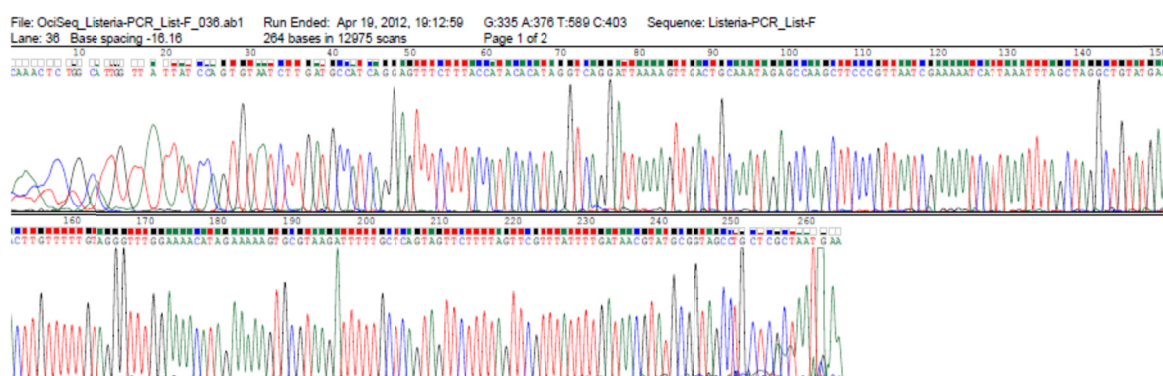
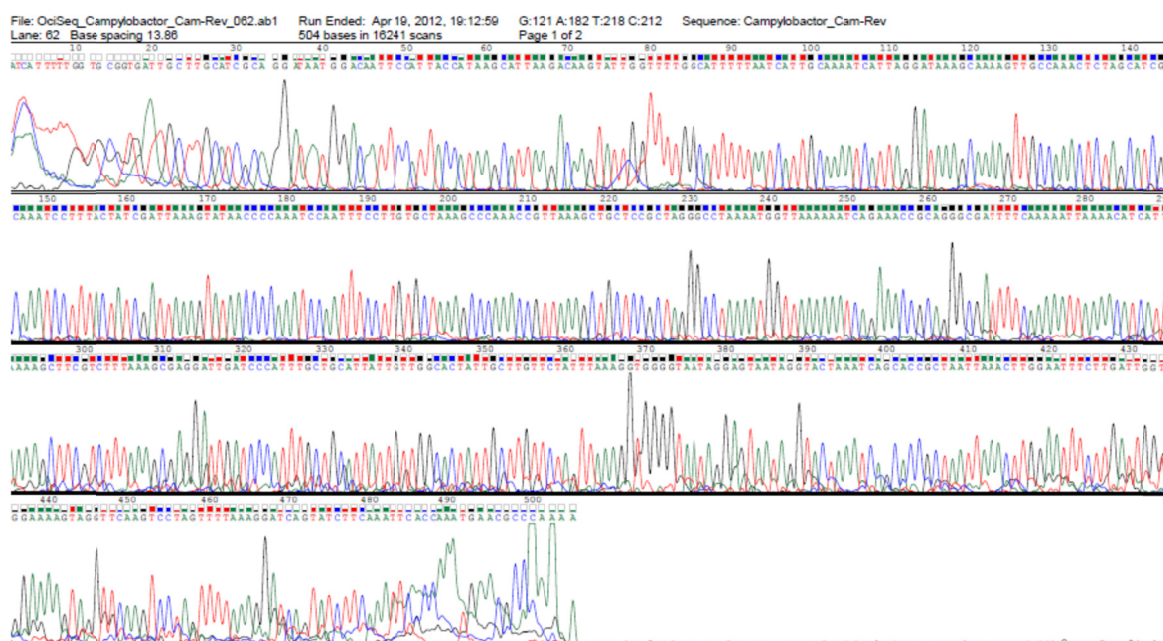


Figure 5. Electropherogram showing amplified DNA sequence data of *Hyp* gene of *Campylobacter jejuni* (Forward primer)



3.6 Screening of Chicken Meat Samples by Multiplex PCR (m-PCR)

In this study, screening of chicken meat samples from retail outlets were carried out to assess the usefulness of the developed m-PCR technique. All the samples screened were not positive either for *Campylobacter jejuni* and *Listeria monocytogenes*. The negative samples were further checked by culture methods and good correlation between these two methods was observed. Rahimi et al. (2010) found higher prevalence of *Campylobacter* (61.7%) in chicken meat from retail markets of Iran. Similarly Kozacinski et al. (2006) conducted a study to assess the microbial quality of chicken meat in Croatian market and reported *L.monocytogenes* in chicken breast and skin (4.76%) but *Campylobacter jejuni* were absent in analysed samples. Similar findings reported by Ahmed and Nashwa (2010) who detected *L. monocytogenes* in chicken meat (16%). Hence, the m-PCR technique developed in this study can be used as a rapid screening test for detection of *Campylobacter jejuni* and *Listeria monocytogenes* from chicken meat within 24 hours.

4. Conclusion

DNA extraction by Phenol Chloroform method was good in purity as well as concentration and cost effective. The developed m-PCR technique could able to detect *Campylobacter jejuni* and *Listeria monocytogenes* from chicken meat within 24 hours even with minimum concentration level of 0.2 ng/μl and 1.0 ng/μl of DNA in a reaction mixture (25 μl). Hence it can be used as rapid screening test.

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