

EXPLORING THE MULTIPLEX PCR FOR DETECTION OF ANIMAL-DERIVED INGREDIENTS IN VEGETARIAN FOODS

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ABSTRACT

Background and objective: The identification of animal-derived ingredients in a vegetarian product is an important issue to protect the consumer from undesirable adulteration, especially, for religious and health reasons. Therefore, it is necessary to establish a rapid method for detecting meat ingredients in processed vegetarian foods. **Materials and methods:** The multiplex PCR assay, based on the amplification of mitochondria *cytochrome b* gene, specific to most of the animals, and *chloroplast DNA*, specific to plant species and served as internal control, were established. Experimentally, a total of 200 vegetarian samples, collected from local markets, were enrolled in the current study. **Results and Conclusion:** 68/200 (counting for 34.0%) samples were positive, which meant having animal-derived ingredients in their foods processing. Also, representative PCR products were conducted sequencing, that showed the animal-derived ingredients. In summary, the results indicated the multiplex PCR could be a reliable method to detect animal-derived ingredients in vegetarian foods applied on a large number of samples in the reality. **Conflict of interest:** The authors declare no conflict of interest.

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Introduction

Most Scholars believe that the effectiveness of health education and behavior change programs depends largely on the use of models and theories of health education and various factors, such as migration, ethnical-racial differences, household income [1,2]. In Vietnam, most of the people are vegetarian eating due to religious, habits, and paying much attention to the issue of their nutrition and health. Therefore, the identification of animal-derived ingredients in a vegetarian product is an important issue to protect the consumer from undesirable adulteration, especially, for religious and health reasons. For this purpose, it is necessary to develop the methods for the detection of animal species in vegetarian foods, which meant that choosing vegetarian food without any animal-derived ingredients. Up to date, numerous analytical methods have been developed based on protein and DNA analysis, such as the protein fractions based technique, or immunological technique, which was carried on the identification of animal tissues in meat products [3-5]. DNA barcodes or probes might be used to identify numerous unknown species [6]. Unfortunately, these methods are often not suitable for the detection of animal-derived ingredients on vegetarian food that are time-consuming and expensive, especially, the complex ingredients.

Advanced DNA technology has led to the rapid development of alternative approaches for species identified. Particularly, polymerase chain reaction (PCR) assay and DNA sequencing were proved to be alternatives to DNA hybridization and protein-based methods because of their accuracy, simplicity, specificity, and sensitivity [7-9]. The current study developed a method for identification of animal-derived species on vegetarian food using PCR assay based on the *cytochrome b* (*cytb*) gene, which is located on the mitochondrial DNA taking more advantages in the species identification, taxonomic and phylogenetic studies [10-13].

Materials and Methods

Samples collection and DNA extraction

In this study, a total of 200 vegetarian products, raw meat (pork and chicken), plant (cabbage) were collected from local markets and supermarkets. The total of DNA was extracted from 1 g of samples using Phenol (pH = 8)/Chloroform methods.

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The quality of DNA was checked on the quantitation by evaluating the absorbance at OD₂₆₀ and OD₂₈₀. The pure preparation of DNA with the OD₂₆₀/OD₂₈₀ ratio of 1.8–2.0 was used to further PCR assay.

Simplex and multiplex PCR assay

In the preliminary phase of this development, the universal primer for *cytb* gene amplification was Cyt-L, Cyto-R (Table 1), as described by Kocher *et al.* (1989)[7], was assessed with DNA extracted from vegetarian foods, and pork meat served as the positive control. Additionally, the other PCR assay, *chloroplast DNA (cpDNA)*, which was specific to plant species, was used as the internal control. The internal control *cpDNA* was amplified using primer according to Taberlet *et al.* (1991)[14] (Table 1).

The PCR was carried out in a total of 15 µL containing 250 ng DNA template, 0.75 unit *iTaq* polymerase (Biorad). PCR reaction was subjected to initial incubation at 95°C for 5 min, followed by 40 cycles at 95°C for 30 s, x°C for 30 s, 72°C for 30 s, and 72°C for 5 min for the final incubation. (Note: x was the annealing temperature for each specific primers to candidate genes, shown in Table 1). Each PCR product was directly loaded onto a 2.0% agarose gel, stained with ethidium bromide, and directly visualized under ultraviolet illumination. Then, the PCR products were sequenced to confirm the specificity of primers. A one-step multiplex PCR was developed using the simultaneous detection of both *cytb* and *cpDNA* to identify the precise, positive of animal-derived ingredients in vegetarian products. The thermal cycling was programmed following the same previously described procedure used in simplex PCR, however, the annealing temperature is 52°C. The results were also observed by electrophoresis on 2.0% agarose.

Results

Simplex PCR specificity and DNA sequencing

In our preliminary phrase of the detection, simplex PCRs were carried out on DNA extracted from raw meat (pork and chicken), plant (cabbage), and vegetarian foods. The results showed that the primer Cyt-L, Cyt-R amplified fragment of 358 bps for both raw meat (Figure 1A, land 5, 6), and one vegetarian sample (Figure 1A, land 2). Meanwhile, no band was observed in plant samples (Figure 1A, land 1), and two vegetarian samples (Figure 1A, land 3, 4). Also, the internal control, *cpDNA*, specific to plant species, was used as an indicator of evaluation of DNA extraction, quality of samples, and quality of PCR. The 426 bps-length bands were observed in the only case of a plant (Figure 1B, land 1), and vegetarian food (Figure 1B, land 3). Contrary to the positive results, in the case of raw meat, no amplifier result was observed (Figure 1B, land 2).

Confirmation of the primers' specificity, 358 bps-length (yielded by Cyt-L, Cyt-R primer), and 426 bps-length (yielded by Clo-F, Clo-R primer) were sequencing, shown in Figure 2. According to results, the signals of peaks in PCR product sequencing were good for nucleotide reading. The results of the BLAST of both sequencings were summarized in Table 2. The amplified sequences, yielded by Clo-F, Clo-R primer, and Clo-F, Clo-R primer, generated in our study revealed to *Gallus gallus* and *Brasica oleracea*, respectively.

Multiplex PCR applied in local vegetarian samples

The multiplex PCR assay was carried out on analogous samples, the set of primers retained the same specificity. Regarding commercial vegetarian foods, collected from local market and supermarket, has been analyzed by multiplex PCR with both Cyt-L, Cyt-R, and Clo-F, Clo-R primer sets, indicated in Figure 3. As for the vegetarian foods, two distinguished bands within 426 bps-length and 358 bps-length (Figure 3, land 3, 6, 7), and only one band with 426 bps-length (Figure 3, land 4, 8), were detected. Also, only one band revealed to *cpDNA* (Figure 3, land 1,2) and *cytb* (Figure 3, land 9, 10) was observed in plant samples and animal samples, respectively. The results of a multiplex PCR assay for the commercial vegetarian foods showed that detection of *Cytb* was successful in 68 out of 200 samples, counting for 34.0%, meant having animal-derived ingredients in their foods processing.

Discussion

Due to religion, habits, and paying much attention to the issue of their nutrition and health, vegetarian foods are required to not contain the animal-derived ingredient. It is necessary to establish a rapid method for detecting meat ingredients in processed vegetarian foods. Up to date, various methods have been applied to detect animal-derived ingredients in food. Among them, DNA-based assays have been developed so far to overcome the limitation of the protein-based method. The limitation of the protein-based assay is the low sensitivity of the high-temperature processed food, resulting in animal proteins were denatured. In the case of DNA-based assays, the DNA molecule was identified as the thermal-stable, and high specific sequence, so DNA is identified as the suitable target for detecting animal-derived ingredients in the high-temperature processed food. Therefore, in this study, the multiplex PCR was developed for the identification of animal-derived species on vegetarian food using PCR assay based on the target gene - *cytochrome b (cytb)* gene and internal control gene - *cpDNA*.

In this study, a specific *cytb* PCR was developed for the identification of animal-derived ingredients in vegetarian foods. In the current study, *cytb* genes were a target for the identification of meat species by PCR. The *cytb* gene of mitochondrial DNA (mtDNA) has been used in widely species identification, taxonomic and phylogenetic studies [12, 15, 16]. Previous studies showed that the use of mtDNA has various advantages such as mtDNA is presented in thousands of copies per cell, compared to nuclear DNA makes it locus ideal for analysis [17].

The results of this simplex PCR as well as multiplex PCR assay showed good evidence for molecular markers linked to the *cytb* and *cpDNA* identification, according to the detection of animal-derivation and plant tissue. The accuracy of the current protocol was confirmed by DNA sequencing, as the results, the signal of sequencing was unique and good for reading, which was homologous to *Gallus gallus*. Additionally, the *cpDNA* product of 426 bps indicated the quality of the DNA extraction and PCR technique. Therefore, these results confirmed multiplex PCR with both Cyt-L, Cyt-R, and Clo-F, Clo-R primer amplification in the current study could be applied in the detection of animal-derived ingredient in vegetarian foods. For the initial application, we pointed out the positive animal-derived in vegetarian foods reached to 34.0%, by the evaluation of *cytb* presence. In a further study, it was necessary to determine the sensitivity of the current PCR assay used in detecting animal-derived ingredient in vegetarian foods. Additionally, in further refinement, the set of Cyt-L and Cyt-R could be more useful for the identification and quantitative analysis of presence animal-derived ingredient in vegetarian foods by the development of quantitative real-time PCR

Conclusion

The present results confirmed the multiplex PCR described and used in the study allows the detection of animal-derived ingredients in vegetarian foods by the amplification of the *cytb* gene as the specific target from animal species. For evaluation of the accuracy, the amplification of *cpDNA*, specific to plant species, was used as the internal control. In our initial application on vegetarian food collected from the local market and supermarket, showed the positive rate was 34.0%. For further study, the sensitivity of the current method may have been studied in the qualitative detection of the target gene.

Conflict of interest

The authors declared that they have no competing interests.

References

1. Claeys E, Uytterhaegen L, Buts B, Demeyer D. Quantification of beef myofibrillar proteins by SDS-PAGE. *Meat Sci.* 1995; 39(2): 177-93. Doi: 10.1016/0309-1740(94)p1819-h.
2. Azizi M, Yazdanpanah A, Hessam S. Investigating the role of Afghan migrants on the health of Shiraz city in 2018. *J. Adv. Pharm. Educ. Res.* 2019;9(S2):114-119.
3. Alshehri MA. Identification of Algae Species Using Advanced Molecular Techniques. *Int. J. Pharm. Res. Allied Sci.* 2020, 9(1):142-159
4. Hsieh YH, Sheu SC, Bridgman RC. Development of a monoclonal antibody specific to cooked mammalian meats. *J Food Prot.* 1998; 61(4): 476-81. Doi: 10.4315/0362-028x-61.4.476.
5. Hsieh YH, Ofori JA. Detection of horse meat contamination in raw and heat-processed meat products. *J Agric Food Chem.* 2014; 62(52): 12536-44. doi: 10.1021/jf504032j.
6. Jamshidi S, Beigrezaei S, Faraji H. A Review of Probable Effects of Antioxidants on DNA Damage. *Int. J. Pharm. Phytopharm. Res.* 2018 Oct 1;8(5):72-9.
7. Kesmen Z, Sahin F, Yetim H. PCR assay for the identification of animal species in cooked sausages. *Meat Sci.* 2007; 77(4):649-53. doi: 10.1016/j.meatsci.2007.05.018.
8. Yang L, Tan Z, Wang D, Xue L, Guan MX, Huang T, Li R. Species identification through mitochondrial rRNA genetic analysis. *Sci Rep.* 2014; 4:4089. doi: 10.1038/srep04089.
9. Mazaheri Nezhad Fard R, Sayyed Asgari F, Ashrafi I, Gharibi S, Khani M. Comparison of Multiple Tube Fermentation and Polymerase Chain Reaction Methods for the Detection of Coliforms in Freshwater. *Appl Food Biotechnol.* 2019; 6(2), 111-117. Doi: 10.22037/afb.v6i2.22484
10. Kocher TD, Thomas WK, Meyer A, Edwards S V, Pääbo S, Villablanca F X, Wilson AC. Dynamics of mitochondrial DNA evolution in animals: amplification and sequencing with conserved primers. *Proceedings of the National Academy of Sciences of the United States of America.* 1989, 86, 6196-6200. Doi: 10.1073/pnas.86.16.6196.
11. Irwin DM, Kocher TD, Wilson AC. Evolution of the cytochrome b gene of mammals. *J Mol Evol.* 1991, 32, 128-144. Doi: 10.1007/BF02515385.
12. Parson W, Pegoraro K, Niederstätter H, Föger M, Steinlechner M. Species identification by means of the cytochrome b gene. *Int J Legal Med.* 2000, 114, 23-28. Doi: 10.1007/s004140000134.
13. Mane BG, Mendiratta SK, Tiwari AK, Narayan R. Sequence analysis of mitochondrial 16S rRNA gene to identify meat species. *J. Appl. Anim.* 2013, 41, 77-81. Doi: 10.1080/09712119.2012.738213.
14. Taberlet P, Gielly L, Pautou G, Bouvet J. Universal primers for amplification of three non-coding regions of chloroplast DNA. *Plant Mol. Biol.* 1991, 17, 1105-1109. Doi: 10.1007/BF00037152.
15. Hsieh HM, Chiang HL, Tsai LC, Lai SY, Huang NE, Linacre A, Lee JC. Cytochrome b gene for species identification of the conservation animals. *Forensic Sci. Int.* 2001, 122, 7-18. Doi: 10.1016/s0379-0738(01)00403-0.
16. Silva-Neto AA, Ferreira PB, Torres RA, Texeira RH, Duarte JM, Barbosa AC, Vargas RC, Garcia JE. Diagnostic Cytochrome b gene profiles for the identification of paca (*Cuniculus paca*) bushmeat: implications for the monitoring of illegal hunting and wildlife trade. *Braz. J. Biol.* 2016, 76, 55-58. Doi: 10.1590/1519-6984.10814.

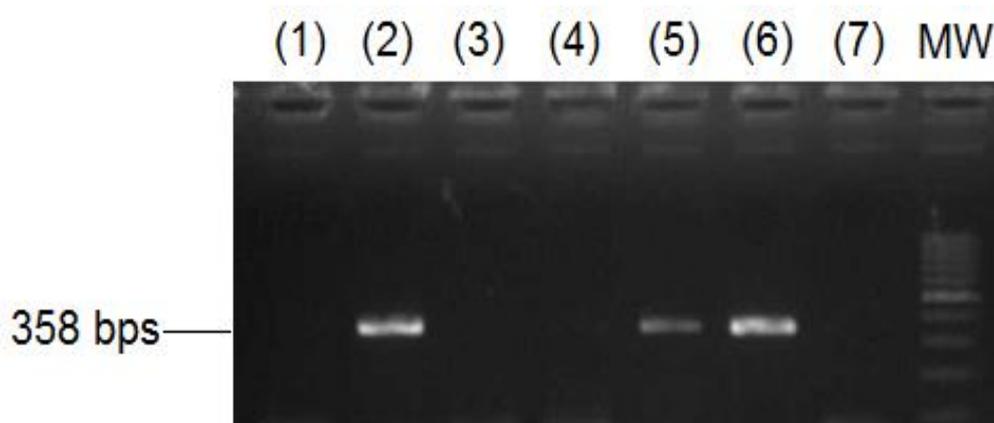
17. Jeihooni AK, Karimi S, Kashfi SH, Mansourian M, Afzali P. Effectiveness of educational program based on Trans-Theoretical model in prevention of osteoporosis in women. *J. Adv. Pharm. Educ. Res.* 2018;8(S2):193-199.

Table 1. Primers' sequences were used in the current study

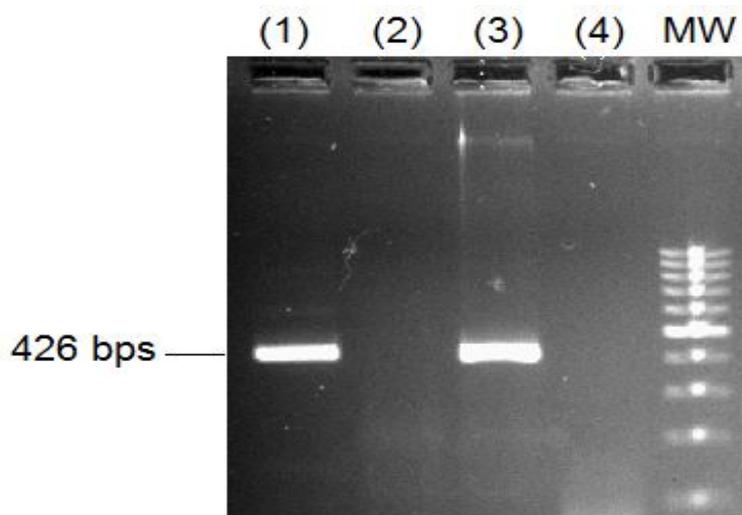
Primer name	Sequence (5' – 3')	x°C
Cyt-L	CCATCCAACATCTCAGCATGATGAAA	52°C
Cyt-R	CCCCTCAGAATGATATTTGTCCTCA	
CLO-F	GGTTCAAGTCCCTCTATCCC	53°C
CLO-R	ATTTGAACTGGTGACACGAG	

Table 2. BLAST results of Sequencing of amplifiers

Sequencing samples	BLASTn (Genbank)			
	Homology	Accession number	Homology rate	E-value
Cyt-L, Cyt-R product	<i>Gallus gallus</i>	DQ512917	97.0%	1e-123
Clo-F, Clo-R product	<i>Brassica oleracea</i>	GQ268028	99.0%	0.0



(A)



(B)

Figure 1. (A) The simplex PCR of DNA with Cyt-L, Cyt-R primer. (1) Plant (cabbage), (2)-(4) Vegetarian food, (5) Raw meat, (6) Raw chicken, (7) Negative control. MW: molecular weight – 100bps. (B) The simplex PCR of DNA with Clo-F, Clo-R primer. (1) Plant species (cabbage), (2) Raw meat (Pork), (3) Vegetarian food, (4) Negative control

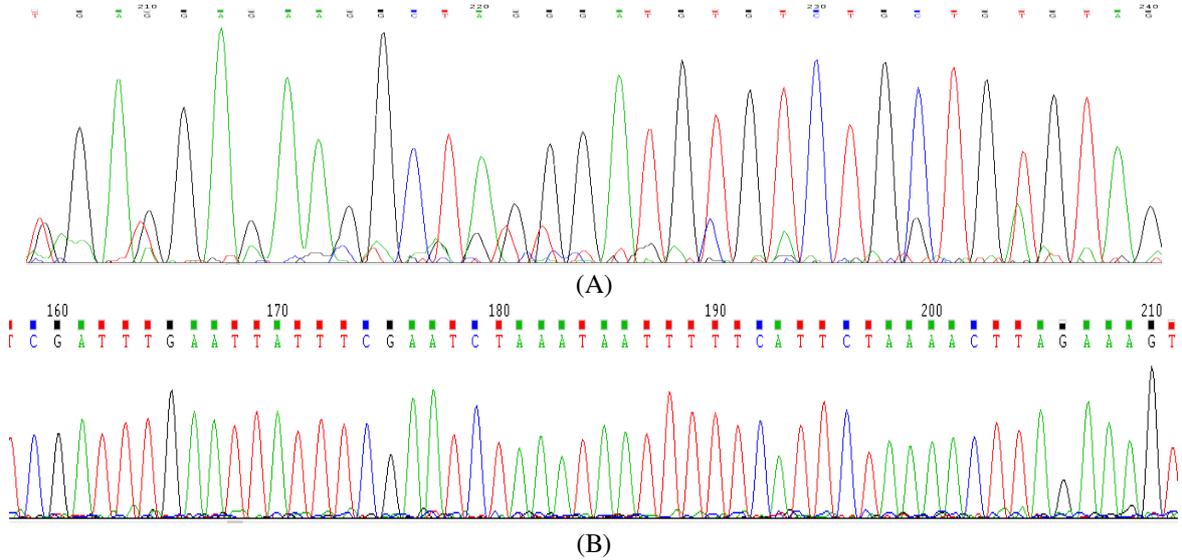


Figure 2. The sequencing of DNA amplifiers with (A) Cyt-L, Cyt-R primer, (B) Clo-F, Clo-R primer

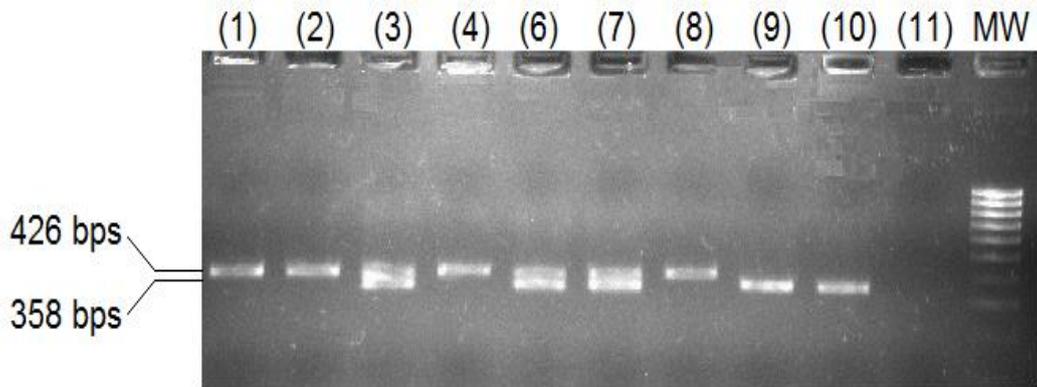


Figure 3. Multiplex PCR on (1) (2) plant samples, (3)-(8) represented commercial vegetarian foods, (9) (10) raw meat, (11) negative control, MW: molecular weight – 100 bps