



**COMPARATIVE DNA EXTRACTION METHOD FROM ADHESIVE TAPE
EVIDENCE FOR PERSON IDENTIFICATION**

By

Natthalak Pakdeenarong

A Thesis Submitted in Partial Fulfillment of the Requirements for the Degree

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Program of Forensic Science

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การเปรียบเทียบวิธีการสกัดสารพันธุกรรมแบบต่าง ๆ ในวัตถุดิบประเภทเห็ดปลวกเพื่อใช้พิสูจน์
เอกลักษณ์บุคคล

โดย

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The Graduate School, Silpakorn University has approved and accredited the Thesis title of “Comparative DNA Extraction Method from Adhesive Tape Evidence for Person Identification” submitted by Miss Natthalak Pakdeenarong as a partial fulfillment of the requirements for the degree of Master of Science in Forensic Science.

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Electrical adhesive tape always found in the crime scene, especially the explosion scene where use electrical adhesive tape for explosive assembling. User's epithelial cells always leaving on the electrical adhesive tape, and can use for DNA extraction to apply in the field of person identification. One of the most important process is DNA extraction from several evidences, this research wants to study and compare the quantity of DNA extraction method from latent fingerprints on electrical adhesive tape evidence from ten volunteers for person identification. The volunteer's touch the three trademarks of electrical adhesive tape which selling in the market with their fingers, Scotch® 3M 1710, Scotch® 3M Super 33+ and YAZAKI. Then, DNA extraction was performed by DNA IQ™ System, Chelex® 100 and Phenol: Chloroform Kit. The extracted DNA samples were quantified by Real-Time PCR to known PCR efficiency and all of data was analyzed by using SPSS program for windows. Chi-square test with the p -value lower than 0.05 was used to indicate statistical significance.

The results of this study indicated that the DNA extraction method were related to the efficiency in PCR process at the 0.05 level of significance ($\chi^2 = 40.833$; $df = 2$; p -value = 0.000). For Chelex® 100 extraction, extracted DNA which has PCR efficiency was observed less than expected results (Standardized residual = -3.3). While the extraction by DNA IQ™ System and Phenol: Chloroform Kit was observed to have PCR efficiency more than the expected results (Standardized residual = 1.6). DNA extraction method gives the highest amount of DNA obtained from electrical adhesive tape were Phenol: Chloroform Kit and DNA IQ™ System. While the extraction by Chelex® 100 gives the least amount of DNA. For electrical adhesive tape trademarks, the statistical analysis showed no significant differences between each electrical adhesive tape trademark and PCR efficiency.

This research showed that the electrical adhesive tape was valuable evidence in forensic science and method probable for DNA extraction on electrical adhesive tape were Phenol: Chloroform extraction and DNA IQ™ System.

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เทปพันสายไฟ เป็นวัตถุพยานที่พบได้เป็นประจำในคดีระเบิดในพื้นที่ 3 จังหวัดชายแดนภาคใต้ โดยเป็นส่วนหนึ่งของการประกอบระเบิด บนเทปขาวมักพบเศษผิวหนังของผู้ใช้งานเทปขาวนั้น ๆ และสามารถนำมาสกัดสารพันธุกรรมเพื่อนำไปใช้พิสูจน์เอกลักษณ์บุคคล การสกัดสารพันธุกรรม เป็นขั้นตอนหนึ่งที่มีความสำคัญต่อการตรวจสอบสารพันธุกรรมจากวัตถุพยาน งานวิจัยนี้จึงทำการศึกษาเปรียบเทียบปริมาณสารพันธุกรรมที่ได้จากวิธีการสกัดสารพันธุกรรมแบบต่าง ๆ ในวัตถุพยานประเภทเทปขาวจากอาสาสมัครจำนวน 10 คน โดยให้อาสาสมัครใช้นิ้วจิ้มเทปพันสายไฟชนิดที่มีขายอยู่ในท้องตลาด 3 เครื่องหมายการค้า คือ Scotch® 3M 1710 Scotch® 3M Super 33+ และ YAZAKI จากนั้นนำมาสกัดสารพันธุกรรมด้วยวิธี DNA IQ™ System Chelex® 100 และ Phenol: Chloroform Kit วัดปริมาณสารพันธุกรรมด้วยวิธี Real-Time PCR เพื่อให้ทราบถึงความสามารถในการนำไปเพิ่มปริมาณ ทำการวิเคราะห์ข้อมูลด้วยโปรแกรม SPSS for windows และวิเคราะห์ผลทางสถิติโดยใช้ Two-Way Chi-square

ผลการวิจัยพบว่าวิธีการสกัดสารพันธุกรรมมีความสัมพันธ์กับความสามารถในการนำไปเพิ่มปริมาณอย่างมีนัยสำคัญทางสถิติที่ระดับนัยสำคัญ 0.05 ($\chi^2 = 40.833$; $df = 2$; $p\text{-value} = 0.000$) โดยการสกัดด้วย Phenol: Chloroform Kit และ DNA IQ™ System ให้ผลมีความสามารถในการนำไปเพิ่มปริมาณมากกว่าที่คาด (Standardized residual = 1.6) ส่วนการสกัดด้วย Chelex® 100 ให้ผลมีความสามารถในการนำไปเพิ่มปริมาณน้อยกว่าที่คาด (Standardized residual = -3.3) วิธีการสกัดสารพันธุกรรมจากวัตถุพยานประเภทเทปขาวที่ให้ปริมาณสารพันธุกรรมมากที่สุด คือ Phenol: Chloroform Kit และ DNA IQ™ System วิธีการสกัดสารพันธุกรรมจากวัตถุพยานประเภทเทปขาวที่ให้ปริมาณสารพันธุกรรมน้อยที่สุด คือ Chelex® 100 ศึกษาความสัมพันธ์ระหว่างเครื่องหมายการค้ากับความสามารถในการนำไปเพิ่มปริมาณพบว่า เครื่องหมายการค้าไม่มีความสัมพันธ์กับความสามารถในการนำไปเพิ่มปริมาณ

การวิจัยนี้แสดงให้เห็นว่าเทปขาวพันสายไฟเป็นวัตถุพยานที่มีคุณค่าทางนิติวิทยาศาสตร์ และสามารถทำการสกัดสารพันธุกรรมจากวัตถุพยานประเภทเทปขาวพันสายไฟได้จากวิธีการสกัดด้วย Phenol: Chloroform และ DNA IQ™ System อย่างมีประสิทธิภาพ

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LIST OF ABBREVIATIONS

Abbreviation	Term
bp	Base pair
cm	Centrimetre
C _T	Threshold cycle
°C	Degree celcius
DEAE	Diethylaminoethyl
DNA	Deoxyribonucleic acid
EDTA	Ethylenediamine tetraacetic acid
HCl	Hydrochloric acid
IPC	Internal PCR control
kbp	Kilo base pair
LCN	Low copy number
mg	Milligram
MgCl ₂	Magnesium chloride
min	Minute
ml	Millilitre
mM	Millimolar
mtDNA	Mitochondrial DNA
Na ₂ EDTA	Disodium ethylenediamine tetraacetate
ng	Nanogram
PCR	Polymerase chain reaction
pg	Picogram
RFLP	Restriction fragment length polymorphism
rpm	Revolutions per minute
SDS	Sodium dodecyl sulfat
sec	Second
SNPs	Single nucleotide polymorphisms

Abbreviation	Term
STR	Short tandem repeat
TBE	Tris (hydroxymethyl) aminomethan – boric EDTA
TE	Tris (hydroxymethyl) aminomethan – EDTA
VNTR	Variable number tandem repeat
μl	Microlitre

CHAPTER I

INTRODUCTION

1. Background and the importance of the problem

Unrest situation in the 3 deep-south provinces of Thailand is one of the most important problems being solved by the Royal Thai Government. Solving this problem requires co-operations from several government bodies including the Royal Thai Police, Ministry of Defense, Ministry of Interior and Ministry of Justice. Major violence problems are bomb trapping, set-up attack, or face-to-face fighting between terrorists and government officers. These cases have to be investigated to find the facts, then proceeding to seek for the real guilty person consequently.

The explosive means “any substance when touching the heat, force of impact or rub against, can transmute from originally become large amount of gas that cause a lot of pressure and heats”.¹ From studying on the explosion cases of the unrest situation in the 3 deep-south provinces, found that, there are many different methods to assemble the explosives. An important component being used is electrical adhesive tape. It has been used to adjoin the explosive material together with the electronics circuit that makes it workable.

Adhesive tape has been used in forensic science, for analysis as a genetic material from the crime scene. There are many kinds of adhesive tape such as paper tape; plastic tape and cloth tape, which have been found in many typical crime scenes. Especially for the electrical adhesive tapes, that has been found as an explosive component in the explosion scenes, always. These tapes are used for assembling the explosive. Therefore, the epithelial cells of the one who had used the adhesive tapes had always been left on it. Then, we can use the DNA extraction to identify the person who had used it. The DNA analysis for person identification in forensic science has been developed continuously, from detecting of the blood group heredity such as, ABO blood group, MNSs blood group, and Rh blood group.

¹Atthapon Chamsuwanavong et al., Forensic science for investigation I, 4th ed. (Bangkok: TCG Printing, 2003), 270.

ABO blood group is the first blood group system discovered and is most important blood group in biological and medical. The person, who has a kind of antigen in his/her erythrocyte, will not have the antibody for that kind of antigen in his/her plasma or serum. Therefore, we can test to identify all kind of blood groups.

MNSs blood group shows the resemble characters of having two gene loci in the close location. There are MN and Ss locus, which each of them is the co-dominant. Each kind of antigen can be tested by reacting with the same antibody, which are anti M, anti N, anti S and anti s.

Rh blood group have the autosomal dominant hereditary relay, the Rh⁺ character prominent is similar to the Rh⁻ one. The Rh blood group has special properties distinct from the ABO and the other one. It has no naturally occurring antibody. The person who has the Rh⁻ will not has anti Rh in his/her body, if he/she has never been immunized by receiving the Rh⁺ blood inside the body before.²

Later on, it had been found that the blood group heredity detecting techniques have it own limitation on discrimination power. It requires many evidence samples to react between antigen and antibody. Therefore, the genetic material has been used in the forensic science from then up to now.

The genetic material or DNA, the shorten form of Deoxyribonucleic acid, is the material derived from all kind of living creatures. It is exist in nucleus and mitochondria of the cells such as blood cells, skin cells, buccal cells, and hair roots. The DNA stores the hereditary information for reproducing of molecular material such as all kinds of proteins both in high-level and low-level life forms. Further than that, it also controls the growth of cells, division of cells, and changing of tissues. Therefore, the DNA of high and low level life forms is the code or blueprint of construction.

Forensic DNA typing for person identification was published in a Nature journal in 1985 by Alex J. Jeffreys. It has been developing continuously until it becomes to the variety of readymade kits now. The most important process to collect genetic information from the evidences is the DNA extraction from several forms of evidences that cannot be expected. This makes the

² Wijarn panich et al., Human genetics, 2nd ed. (Bangkok : Pikanes printing, 2004), 271-272, 288, 292.

difficulty in DNA typing. Therefore, the theories to be used for DNA extraction from the evidences must be the most appropriate. There are many chemical extraction methods and procedures for getting the most DNA quantity from evidences, without the contamination caused failure in DNA typing.

For the reason aforementioned, the researchers want to study and to compare the quantity of DNA extracted from various methods for person identification purpose, which are collected from the latent fingerprints on electrical adhesive tape evidence.

2. Research objectives

- 2.1 To determine the relationships among DNA extraction methods and its efficiency in PCR processes, for the latent fingerprints left on electrical adhesive tape evidence.
- 2.2 To determine the relationships among various brands of electrical adhesive tape and its efficiency in PCR processes.
- 2.3 To study problems and to develop recommendations, for the DNA extraction collected from latent fingerprints left on the electrical adhesive tape evidence.

3. Research hypotheses

- 3.1 The DNA extraction methods relate to the efficiency in PCR process.
- 3.2 The trademarks of electrical adhesive tape relate to the efficiency in PCR process.

4. Scope of research

To study the quantity of extracted DNA from ten volunteers by using nanogram per microlitre (ng/ μ l) as analyse unit, by controlling of the volunteers' finger touching on the three trademarks of electrical adhesive tapes available in the market, for three samples per one trademark, per volunteer.

5. Conceptual framework

A conceptual framework for the research, which provides a guideline for conducting the research. The details are illustrated in Figure 1

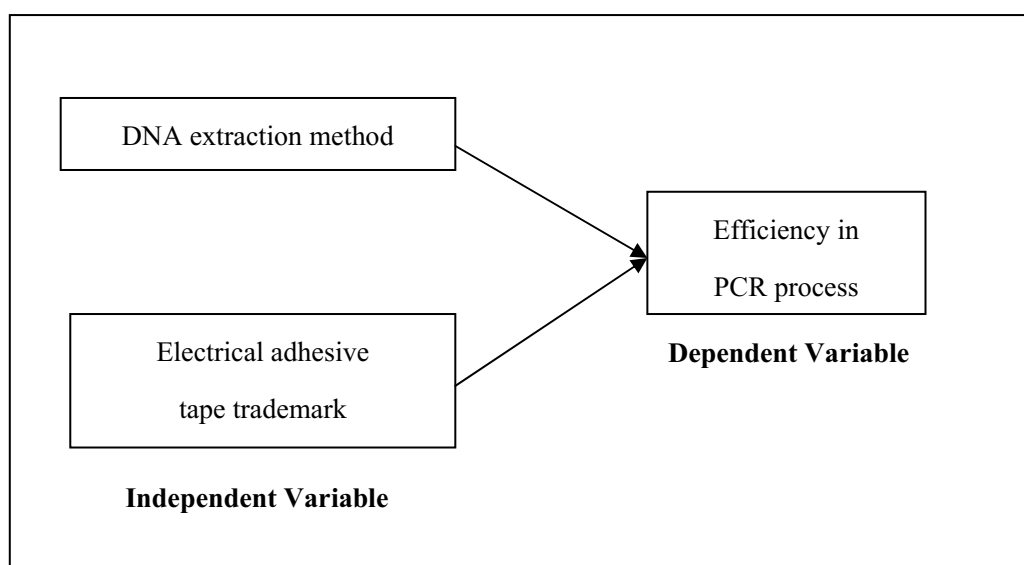


Figure 1 A conceptual framework for the research.

6. Research usefulness

- 6.1 To know the effective methodology guidelines of DNA extraction from electrical adhesive tape evidences.
- 6.2 To know the problems and recommendations about DNA extractions from the latent fingerprints left on electrical adhesive tape evidences.
- 6.3 To use the DNA profiles that extraction from electrical adhesive tapes to be DNA database and link to the suspect for solve the unrest situation in the 3 deep-south provinces of Thailand.

CHAPTER II

LITERATURE REVIEW

Forensic science combines and applies every fields of science altogether in favor to the justice processes. Therefore, the scopes of forensic science are wide spread, starting from proving of a small case of stealing up to the serious crime cases, such as rape, sexual assault and homicide. However, the forensic scientists should certainly aim to use the science knowledge to prove and to present the analysis result clearly and transparently. Forensic sciences comprise of many fields such as Forensic Odontology, Forensic Osteology, Forensic Pathology, Forensic Entomology, Forensic Toxicology, Forensic Medicine and Forensic DNA typing.

‘DNA fingerprinting’ or DNA typing (profiling) was first described in 1985 by an English geneticist named Alec Jeffreys. By developing a technique to examine the length variation of these DNA repeat sequences, Dr. Jeffreys created the ability to perform human identity tests.¹ The applications of human identity are crime solving, accident victims, soldier in war, paternity testing, inheritance claims, missing persons investigations, and convicted felons database.

DNA is present in every nucleated cell and is therefore present in biological materials left at crime scenes. It has been successfully isolated and analyzed from a variety of biological materials,² For example: buccal cells, blood, tissues, bones, hair roots and epithelial cells.

Some epithelial cells always leavings on the object, including the floor, pen, table, glass, clothes and electrical adhesive tape, was used to DNA extraction for apply in the field of person identification.

Electrical adhesive tapes were used in many ways. Therefore, it has always been found in the crime scenes, especially for the explosion scenes in the unrest situation of three deep-south provinces of Thailand. The terrorists have used the electrical adhesive tape to assemble the explosives. These explosions cause varied damage to the country including international

¹J.M. Butler, Forensic DNA Typing: Biology and Technology behind STR markers, (Harcourt Place London : Academic Press, 2001), 2-3.

²Ibid., 171.

relationship, economy, politics, social, religion, personnel and education. So, there are many set-ups of co-operations between several government bodies including the Royal Thai Police, Ministry of Defense, Ministry of Interior, and Ministry of Justice, for solving the unrest.

1. Explosives and explosions

An explosive is a material capable of rapid conversion from either a solid or liquid to a gas with resultant heat, pressure, and loud noise.³ The effects of an explosion can all be explained by understanding what happens when an explosion or detonation takes place. Solid or liquid fuels combine with oxygen to form gaseous products such as carbon dioxide and other products that are converted to gases from the heat of the combustion. These very hot gases expand rapidly away from the origin of the explosion (the bomb seat). These rapidly moving gases create three primary effects: blast pressure, fragmentation, and thermal or heat effects.⁴

Collecting physical evidence at the scene consists of the search for and recovery of items that may lead to information about the nature and type of explosive and the identity of the suspect. The area should be searched for the fusing mechanism of the bomb. Items such as timing mechanisms, batteries, pieces of wire, safety fuse, blasting cap debris, and the like may yield information about the way in which the bomb was set to detonate.

The investigator should not forget to search for other evidence besides the bomb debris. Items such as fingerprints, tire tracks, tool marks, electrical adhesive tape, and the like are valuable and must not be overlooked.⁵

2. Adhesive tape

Electrical, adhesive, masking, and cellophane tape are sometimes recovered at crime scenes. The tape may have been used to bind a victim or to tape two objects together. If a roll of tape is

³Barry A.J. Fisher, Techniques of crime scene investigation, 5th ed. (Florida : CRC Press, 2000), 315.

⁴Max M. Houck and Jay A. Siegel, Fundamental of Forensic Science, (Oxford : Elsevier Limited, 2006), 492-493.

⁵Barry A.J. Fisher, Techniques of crime scene investigation, 327-328.

located in the suspect's belongings, it may be possible to piece together the portion of tape from the scene and that found in the suspect's possession. If the tear at the end of the tape is ragged enough, a conclusive statement about a common source may be made. If the cut on the tape is very sharp, such as may be made by scissors, only a statement about class characteristics can be made.⁶

Adhesive tape has been used in forensic science, for analysis as a genetic material from the crime scene. Especially for the electrical adhesive tapes, that has been found as an explosive component in the explosion scenes, always. These tapes are used for assembling the explosive. Therefore, the epithelial cells of the one who had used the adhesive tapes had always been left on it. Then, we can use the DNA extraction to identify the person who had used it.

The electrical adhesive tape is very sticky, an adhesive tape comprising of a layer of adhesive material, a cured butyl rubber based composition formed by compounding a portion of a butyl rubber copolymer.⁷ Therefore, extraction of the DNA on electrical adhesive tape is difficult.

3. Epithelial cells

Epithelium is a tissue composed of a layer of cells. Structurally, epithelium lines the outside (skin) can be divided into two regions, the outer epidermis and the thicker inner dermis. The epidermis is composed of several layers of cells that are often described as being a keratinized stratified squamous epithelium.⁸

Functions of epithelial cells include protection, homeostasis, excretion, temperature regulation, vitamin D production, sensory perception, psychosocial function, and wound healing.⁹

⁶Barry A.J. Fisher, Techniques of crime scene investigation, 5th ed. (Florida : CRC Press, 2000), 212.

⁷Adhesive tape compositions. [Online], accessed 25 September 2008. Available from <http://www.patentstorm.us/patents/4588637.html>

⁸Stephen R.B. et al., Cell Biology (A Short Course). 2nd ed. (New Jersey: John Wiley & Sons; 2004).

⁹Christina Marino, Skin Physiology, Irritants, Dry Skin and Moisturizers [Online], accessed 12 September 2008. Available from <http://www.lni.wa.gov/Safety/Research/Dermatitis/> 1-888-66-SHARP

Epithelial cells can be deposited when an object is touched.¹⁰ The amount of cellular material transferred depends upon the amount of time the skin is in contact with the object; the amount of pressure applied; and the presence of fluid such as sweat to mediate the transfer. Some people transfer their skin cells more readily than others; these people are classified as good shedders.¹¹ In most cases the number of cells is very low and the success rate of DNA profiling is limited. Screening methods, for example using the reagent ninhydrin, which detects the presence of amino acids, can be helpful in identifying samples that are likely to contain epithelial cells.¹²

4. Biological evidence collection

The methods used for collection will vary depending on the type of sample. Dry stains and contact marks on large immovable items are normally collected using a sterile swab that has been moistened with distilled water.¹³ Lifting from the surface using high quality adhesive tape is an alternative method for collecting epithelial cells.¹⁴ Liquid blood can also be applied to FTA[®] paper that is impregnated with chemicals to prevent the action of microbial agents and stabilize the DNA.

Smaller moveable objects, such as weapons, which might contain biological material are packaged at the scene of crime and examined in the controlled environment of the forensic laboratory. The same range of swabbing, scraping and lifting techniques as used in the field can

¹⁰van Oorschot, R.A.H., and M.K. Jones, "DNA fingerprints from fingerprints," Nature (1997) : 387,767–767.

¹¹G.N. Ruttly et al., "The effectiveness of protective clothing in the reduction of potential DNA contamination of the scene of crime," International Journal of Legal Medicine (2003) : 117, 170–174.

¹²K. Anslinger et al., "Ninhydrin treatment as a screening method for the suitability of swabs taken from contact stains for DNA analysis," International Journal of Legal Medicine (2004) : 118, 122–124.

¹³M. Benecke, "Forensic DNA samples—collection and handling," Encyclopedia of Medical Genomics and Proteomics (2005) : 500–504.

¹⁴D. Hall and M. Fairley, "A single approach to the recovery of DNA and firearm discharge residue evidence," Science and Justice (2004) : 15–19.

be employed to collect the biological material.¹⁵ Clothing taken from suspects and victims presents an important source of biological evidence. This is also analysed in the forensic biology laboratory where stains and contact areas can be recorded and then cut out or swabbed.

5. DNA

5.1 Nuclear DNA

Nuclear DNA is located in the nucleus and found in all body cells, except for red blood cells. Each person has the same DNA in every part of the body, including buccal cells, blood, tissues, and bones. Each person receives half of his or her DNA from his/her father and another half from his/her mother. The DNA of each person is unique to that person, except for identical twins, whose DNA are exactly the same.¹⁶⁻¹⁷

5.2 Mitochondrial DNA (mtDNA)

Mitochondrial DNA is found in the mitochondrial and is passed from a mother to her children. In short, mtDNA shows maternal lineage.¹⁸

6. DNA extraction

DNA can be extracted from a wide range of samples such as whole blood, blood stains, semen, hair roots, bones, saliva and skin epithelial cells. The DNA extraction can be disruption of the cellular membranes, resulting in cell lysis, protein denaturation, and the separation of DNA from the denatured protein and other cellular components. There are many methods available for extracting DNA. The choice of which method to use depends on a number of factors, including

¹⁵L. Kobilinsky, Liotti T.F., and Oeser-Sweat J., DNA: Forensic and Legal Applications, (New Jersey : John Wiley & Sons, 2005).

¹⁶Langford A. et al., Practical Skills in Forensic Science, (Pearson Practice Hall, 2005).

¹⁷J.M. Butler, Forensic DNA Typing: Biology and Technology behind STR markers, (Harcourt Place London : Academic Press, 2001), 17-20.

¹⁸Chulalongkorn University and Royal Thai Police, National guideline of forensic science services for disaster victim identification in Thailand, (Bangkok : Chulalongkorn University, 2007), 185.

the sample type and quantity. Different extraction methods are used to isolate DNA from each different sample.

6.1 Conventional DNA extraction methods

6.1.1 Organic (Phenol-Chloroform) extraction. The phenol–chloroform method has been widely used in molecular biology. Organic extraction involves the serial addition of several chemicals. Sodium dodecylsulfate (SDS) and proteinase K are added to break open the cell walls and to break down the proteins. Next a phenol/chloroform mixture is added to separate the proteins from the DNA. The DNA is more soluble in the aqueous portion of the organic–aqueous mixture. When centrifuged, the unwanted proteins and cellular debris are separated away from the aqueous phase and double stranded DNA molecules can be cleanly transferred for analysis. The method produces clean DNA but has some drawbacks: in addition to the toxic nature of phenol, multiple tube changes are required and the process is labour intensive.¹⁹⁻²⁰

6.1.2 Inorganic methods. Inorganic methods are based on the use of either sodium chloride or lithium chloride to ‘salt out’ proteins from mixture of DNA and protein. Procedures using salt have been largely used to extract DNA from blood, cytological samples, and soils material, and proved to be less laborious and toxic than the phenol-chloroform technique while the salting-out extraction method is as efficient as the phenol-chloroform extraction.

6.1.3 Chelex[®] extraction. The Chelex[®] 100 method was one of the first extraction techniques adopted by the forensic community. Chelex[®] 100 is a resin that is composed of styrene-divinylbenzene copolymers containing paired iminodiacetate ions.²¹ The resin has a very high affinity for polyvalent metal ions, such as magnesium (Mg^{2+}); it chelates the polyvalent metal ions and effectively removes them from solution. The extraction procedure is very simple. In most protocols, biological samples such as bloodstains are added to a 5% Chelex suspension and boiled for several minutes to break open the cells and release the DNA.

¹⁹J.M. Butler, Forensic DNA Typing: Biology and Technology behind STR markers, (Harcourt Place London : Academic Press, 2001), 44.

²⁰William Goodwin, Adrian Linacre and Sibte Hadi, An Introduction to Forensic Genetics, (England : John Wiley & Sons Ltd, 2007), 29-30.

²¹P.S. Walsh et al., “Chelex-100 as a medium for simple extraction of DNA for PCR-based typing from forensic material,” Biotechniques 10 (1991) : 506–513.

The cellular material is incubated with the Chelex[®] 100 suspension at 56 °C for up to 30 minutes. Proteinase K is often added at this point. This incubation is followed by 8–10 minutes at 100 °C to ensure that all the cells have ruptured and that the protein is denatured. The tube is then simply centrifuged to pellet the Chelex[®] 100 resin and the denatured protein at the bottom of the tube, leaving the aqueous solution containing the DNA to be used in PCR.

The major advantages of this method are: it is quick, simple, the cost is very low, avoids the use of harmful chemicals and does not involve the movement of liquid between tubes. The DNA extract produced using this method is relatively crude but sufficiently clean in most cases to generate a DNA profile.

6.2 Commercial DNA extraction kits

6.2.1 QIAamp extraction. QIAamp[®] (QIAGEN, CA, USA) DNA extraction kits are widely used for DNA extraction from forensic samples.²² QIAamp membrane is a silica column-based membrane with a DEAE anion-exchange resin that interacts with the negatively charged phosphate of the DNA backbone to remove the DNA from other cellular components after the cells are broken open. Impurities are washed from the column with medium ionic strength buffers. The DNA remains bound until eluted with a high-salt buffer. The clean DNA is collected from the bottom end of the column in a drop of liquid.²³

6.2.2 DNA IQ[™] system. A novel approach to quantification is used in the commercially available DNA IQ[™] Isolation System (Promega Corporation). The isolation method is based on salting-out and binding to silica: a very specific amount of silica coated beads is added to the extraction and these bind a maximum amount of DNA; therefore, when the DNA is eluted from the beads the maximum concentration is known. It has the advantage of combining the extraction and quantification steps but has the disadvantage of not being human specific.²⁴

²²Kathryn Sinclair and Victoria M. McKechnie, “DNA extraction from stamps and envelope flaps using QIAamp and QIAshredder,” *Journal of Forensic Science*. 1 (2000) : 229-230.

²³McGillivray B., “The role of Victorian emergency nurses in the collection and preservation of forensic evidence: a review of the literature,” *Accident and Emergency Nursing* 13, 2 (2005) : 95-100.

²⁴William Goodwin, Adrian Linacre and Sibte Hadi, *An Introduction to Forensic Genetics*, (England : John Wiley & Sons Ltd, 2007), 36.

6.2.3 FTA[®] paper. FTA[®] paper is an absorbent cellulose-based paper that contains four chemical substances to protect DNA molecules from nuclease degradation and preserve the paper from bacterial growth. As a result, DNA FTA[®] paper is stable at room temperature over a period of several years.²⁵

FTA[®] paper as a method for sample collection and storage, particularly from buccal swabs and fresh blood samples. Once a sample is applied to the FTA[®] paper it is stable at room temperature for several years. Cellular material lyses on contact with the FTA[®] paper and the DNA becomes bound to the paper. To analyse the DNA sample, the first step is to take a small region of the card, normally a 2-mm diameter circle, place it into a 1.5-ml tube and the non-DNA components are simply washed off, leaving only DNA on the card. The small circle of FTA[®] paper is then added directly to a PCR. The FTA paper extractions are very simple to perform and do not require multiple tube changes, thus reducing the possibility of sample mixing.²⁶

7. PCR inhibition

Outdoor crimes may leave body fluids such as blood and semen on soil, sand, wood, or leaf litter that contain substances which may co-extract with the perpetrator's DNA and prevent PCR amplification. Textile dyes, leather, and wood from interior crime scenes may also contain DNA polymerase inhibitors.²⁷ Inhibitors can interfere with the cell lysis necessary for DNA extraction, interfere by nucleic acid degradation or capture and inhibit polymerase activity thus preventing enzymatic amplification of the target DNA.

Samples containing PCR inhibitors often produce partial profile results that look similar to a degraded DNA sample. Thus, failure to amplify the larger STR loci for a sample can be either due to degraded DNA where there are not enough intact copies of the DNA template or due to the presence of a sufficient level of PCR inhibitor that reduces the activity of the polymerase.

²⁵J.M. Butler, Forensic DNA Typing: Biology and Technology behind STR markers, (Harcourt Place London : Academic Press, 2001), 45.

²⁶William Goodwin, Adrian Linacre and Sibte Hadi, An Introduction to Forensic Genetics, (England : John Wiley & Sons Ltd, 2007), 29-30.

²⁷Bessetti J., An introduction to PCR inhibition. Promega Corporation. March, (2007).

Reduced size STR amplicons can aid in recovery of information from a sample that is inhibited since smaller PCR products may be amplified more efficiently than larger ones.²⁸

8. Tandem repeats

Two important categories of tandem repeat have been used widely in forensic genetics: minisatellites, also referred to as variable number tandem repeats (VNTRs); and microsatellites, also referred to as short tandem repeats (STRs).

8.1 Variable number tandem repeats – VNTRs

VNTRs were the first polymorphisms used in DNA profiling and they were successfully used in forensic casework for several years.²⁹

Each VNTR contain a define sequence of nucleotides about 9-50 bp long repeated a number of times in a tandem fashion. The longer of the minisatellite can vary from several 100 bp to over 20 kbp. The technique used to examine the VNTRs was called restriction fragment length polymorphism (RFLP).³⁰

The use of VNTRs was, however, limited by the type of sample that could be successfully analysed because a large amount of high molecular weight DNA was required. Interpreting VNTR profiles could also be problematic. Their use in forensic genetics has now been replaced by short tandem repeats (STRs).

8.2 Short tandem repeats – STRs

STRs are currently the most commonly analysed genetic polymorphism in forensic genetics. STR loci are spread throughout the genome including the 22 autosomal chromosomes and the X and Y sex chromosomes. They have a core unit of between 1 and 6 bp and the repeats typically range from 50 to 300 bp. The majority of the loci that are used in forensic genetics are

²⁸J.M. Butler, Forensic DNA Typing: Biology and Technology behind STR markers, (Harcourt Place London : Academic Press, 2001), 150.

²⁹A.J. Jeffreys et al., “Positive identification of an immigration test-case using human DNA fingerprints,” Nature 317 (1985) : 818–819.

³⁰Keiji T. and Alec J.J., “Human tandem repeat sequences in forensic DNA typing” Legal Medicine 7(2005) : 244-250.

tetranucleotide repeats, which have a four base pair repeat motif.

STRs satisfy all the requirements for a forensic marker. They are robust, leading to successful analysis of a wide range of biological material, the results generated in different laboratories are easily compared, highly discriminatory, very sensitive, requiring only a few cells for a successful analysis and there is a large number of STRs throughout the genome that do not appear to be under any selective pressure.³¹

8.3 Single nucleotide polymorphisms – SNPs

The simplest type of polymorphism is the SNP; single base differences in the sequence of the DNA. SNPs are formed when errors (mutations) occur as the cell undergoes DNA replication during meiosis. SNPs normally have just two alleles, for example one allele with a guanine and one with an adenine, and therefore are not highly polymorphic and do not fit with the ideal properties of DNA polymorphisms for forensic analysis. However, SNPs are so abundant throughout the genome that it is theoretically possible to type hundreds of them. This will make the combined power of discrimination very high. It is estimated that to achieve the same discriminatory power that is achieved using 10 STRs, 50 – 80 SNPs would have to be analysed.³² With current technology, this is much more difficult than analysing 10 STR loci.³³

9. Relate articles

In 2002, Schulz and Reichert tried to assess the potential use of latent fingerprints as a DNA source for STR typing. Magnetic powder, soot powder and scotch tape were used for visualization and archiving fingerprints in Germany were tested for their PCR inhibitory characteristics. Fingerprints were placed on clean glass surfaces, visualized and tested for their usefulness as a DNA source. The scotch tape was removed from the evidence cards and fingerprints were cut out

³¹William Goodwin, Adrian Linacre and Sibte Hadi, An Introduction to Forensic Genetics, (England : John Wiley & Sons Ltd, 2007), 12-13.

³²P. Gill, “Anassessment of the utility of single nucleotide polymorphisms (SNPs) for forensic Purposes,” International Journal of Legal Medicine 114 (2001) : 204–210.

³³M. Krawczak, “Informativity assessment for biallelic single nucleotide polymorphisms,” Electrophoresis 20 (1999) : 1676–1681.

from the scotch tape, the sample was cut into small pieces and DNA isolation was carried out using the InViSorbTM Forensic Kit I (InViTek GmbH, Berlin). Obtained DNA was quantified and tested in an STR system at locus FGA. The results showed that 48 scotch tape-archived fingerprints, nine could be successfully amplified and typed. Four of these were visualized with soot and five with magnetic powder. It proved possible to type fingerprints removed from the surface with scotch tape and showed that magnetic powder, soot powder and scotch tape not disturbed DNA amplification.³⁴

In 2003, Pesaresi et al. recovered that DNA can be successfully extracted from fingerprints and analysed using short tandem repeat (STR) profiling. The fingerprints of 11 persons were applied on the following clean substrates, glass, metal and wood. Fingerprints were prepared in two different ways: pressure at a standard time of 30 s, and tangential contact with a rolling friction effect on the skin. In addition, dactyloscopic powders were investigated for their influence on DNA. DNA was extracted with phenol–chloroform, quantified using the dot-blot procedure and concentrated using the Microcon-30 procedure (Amicon, Beverley, MA, USA). Using the AmpF/STR Profiler Plus kit. PCR products were electrophoresed on an ABI Prism 310 Genetic Analyzer. The results show that it is possible to type DNA from biological material left by simple skin contact on several kinds of substrate. Unambiguous profiles were obtained from fingerprints left by subjects with clean hands, and mixed profiles were often found in DNA of fingerprints from subjects with unwashed hands.³⁵

In 2003, latent fingerprint on paper was used in STR genotyping and mtDNA sequencing by Balogh et al. In them work, commercial office printer paper was used as starting material. In order to evaluate the performance of latent fingerprint analysis in a criminal case, experiments with varying conditions were carried out to improve our understanding of low copy number (LCN) DNA typing. After optimising the extraction methods to achieve increased sensitivity, the examination of touched paper can routinely yield the STR profile of the individual who has

³⁴M.M. Schulz and W. Reichert, “Archived or directly swabbed latent fingerprints as a DNA source for STR typing,” *Forensic Science International*, 127 (2002) : 128–130.

³⁵M. Pesaresi et al., “Qualitative and quantitative analysis of DNA recovered from fingerprints,” *International Congress Series*, 1239 (2003) : 947–951.

touched it. In the touching period experiment, paper was touched by the four donors for seven different handling time periods ranging from 1 to 60 s. Complete STR profile indicating that the proportion of successfully detected DNA profiles is not dependent on touching period. The experiment time of day was introduced to determine whether there is a difference between fingerprints prepared in the morning, during the course of a day at noon, after sports and in the evening. The results show that the time of day experiment shows no significant differences between fingerprints deposited in the morning without prior hand washing and during the normal course of a day. For these results, a fingerprint can therefore be considered as a potential source of DNA for genetic identification.³⁶

In 2003, Richard and Howard tried to use hydrophilic adhesive tape for collection of evidence for forensic DNA analysis. In this study, the authors have developed a method that employed a hydrophilic adhesive tape (HAT) for collecting DNA evidence. The HAT method was used to remove surface cells from relatively hairless areas on the body. The area examined were ankle, arm, behind the ear, between fingers and back of the neck. The HAT was then dissolved in the extraction buffer. DNA typing was performed at vWA, TH01, F13A1, and FES loci using the short tandem repeat (STR) analysis. The results show that the samples collected from ear give the best results with a success rate of 100%. All subjects tested by this method had known STR genotypes established from buccal swabs. The authors' results suggest that the HAT method can be used as a less invasive method for collecting biological evidence for forensic DNA analysis. In addition, this collection method should reduce the risk of DNA degradation due to the moisture, which is encountered using conventional collecting methods.³⁷

In 2006, Prinz et al. developed methodologies were tested on control samples as well on fingerprints deposited on a variety of substrates such as credit cards, keys, and pens. All samples were amplified in triplicate to confirm the presence of each allele and to detect drop-ins. Overall the modifications implemented produced reproducible results for DNA titrated to 20 pg. For DNA

³⁶M. Kinga Balogh et al., "STR genotyping and mtDNA sequencing of latent fingerprint on paper," *Forensic Science International*, 137 (2003) : 188–195.

³⁷Richard C. Li and Howard A. Harris, "Using hydrophilic adhesive tape for collection of evidence for forensic DNA analysis," *Journal of Forensic Sciences*, 48 (2003) : 1–4.

dilutions, 25 pg routinely resulted in full profiles, and 12.5 pg determined 76.9% of the database loci tested. Similarly, for the touched objects, 75.8% of the 20-pg to 100-pg samples yielded database-eligible profiles; the remaining samples either were mixtures or contained an insufficient number of allelic calls. The three-amplification approach was crucial and produced more complete profiles with confidence in the allelic assignments. DNA amounts below 20 pg did show partial profiles with correct allelic determinations that could have been compared in a specific case but were often too incomplete for database entry.³⁸

In 2007, Aree Lempan tried to investigate DNA recovery from forensic clothing samples by tape-lift. The three different DNA extraction methods were Chelex[®] 100, ChargeSwitch[®] Forensic DNA Purification Kit and QIAamp[®] DNA Mini Kit. The results showed that DNA extraction by ChargeSwitch[®] Forensic DNA Purification Kit was the most effective method for extraction of DNA from tape-lifting. Using 3100 Genetic Analyzer (Applied Biosystems) for DNA typing, DNA profiles of the volunteers' epithelial cells were matched to those collected from corresponding clothing samples. These results demonstrated that DNA typing from tape-lift was possible.³⁹

In 2008, Lagoa et al. compared the application of autosomic STR, Y-STR and mini STR markers on fingerprints genetic analysis, increasing the number of PCR cycles as a strategy to accomplish more sensitivity. 180 fingerprints left on slides were arranged in 60 groups of 3. Each group was swabbed with sterile swab moistened and dry swab. DNA was extracted with phenol: chloroform: isoamyl alcohol, and concentrated with Microcon[®] spin columns. Samples were analyzed with AmpF/STR[®] Identifiler[™] (Applied Biosystems) kit, AmpF/STR[®] Yfiler[™] (Applied Biosystems) kit and two mini STR multiplex using, respectively 34, 36 and 36 cycles. PCR products were separated by 3100 AB Prism Genetic Analyzer. The number of detected alleles increased when miniSTR were used; however, contamination level was higher. AmpF/STR[®] Yfiler[™] kit revealed high sensitivity to DNA degradation and it seems less efficient

³⁸ Mechthild Prinz et al., "Maximization of STR DNA typing success for touched objects," *International Congress Series*, 1288 (2006) : 651–653.

³⁹ Aree Lempan, "DNA recovery from forensic clothing samples by tape-lift," (M.Sc. dissertation, Mahidol University, 2007), 37-41.

than AmpF/STR[®] Identifier[™] kit. They concluded that mini STR are the best choice, though strict guidelines and caution are required when interpreting mini STR low copy number (LCN) profiles.⁴⁰

In 2008, Sewell et al. tried to investigate the various factors affecting DNA profiling from DNA recovered from fingerprints deposited on paper before and after fingerprint enhancement treatments. QIAamp DNA mini kit was compared to DNeasy[®] plant mini kit (both from QIAGEN[®], UK) for efficiency in recovering DNA from both saliva and fingerprints on paper. The DNeasy[®] plant mini kit (QIAGEN[®]) was found to improve DNA recovery from paper by over 150% compared with the QIAamp[®] mini kit. Furthermore, this study found that whilst certain paper types, such as newspaper, magazine and filter paper allowed for the good recovery of DNA, common office paper and white card, strongly interfered with the recovery of DNA resulting in poor quality profiles.⁴¹

Based on related literature, a fingerprint can therefore be considered as a potential source of DNA for genetic identification and DNA from fingerprint can be successfully extracted and analysed using short tandem repeat (STR) profiling. In addition, fingerprints deposited is not dependent on touching period and time of day.

Adhesive tape has been used in forensic science, for analysis as a genetic material from the crime scene. Extracted DNA from adhesive tape could be successfully amplified and typed. It proved possible to type fingerprint removed from the surface with adhesive tape and showed that adhesive composition in adhesive tape not disturbed DNA amplification.

⁴⁰ A.M. Lagoa, T. Magalhães and M.F. Pinheiro, “Autosomal STR, Y-STR and miniSTR markers evaluation for genetic analysis of fingerprints,” Forensic Science International: Genetics Supplement Series, (2008) : n.pag.

⁴¹ Jonathan Sewell et al., “Recovery of DNA and fingerprints from touched documents,” Forensic Science International: Genetics, 2 (2008) : 281–285.

CHAPTER III

MATERIALS AND METHODS

1. Supplies

Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape from Tesco Lotus Department Store (Bangkok, Thailand), Scotch[®] 3M 1710 Vinyl Electrical Tape and YAZAKI P.V.C. Tape from a shop (Nontaburi, Thailand), Chelex[®] 100 resin (Bio-RAD, CA, USA), Phenol: Chloroform Kit (PIERCE, CA, USA), Microcon[®] 100 (Millipore, MA, USA). Chemicals and reagents supplied from Promega Corporation (Medison, WI, USA) as: DNA IQ™ System, Proteinase K, TE buffer and Hypure water. Following Supplies were obtained from Applied Biosystems (Foster City, CA, USA): 10X TBE buffer, Hi-Di formamide, Quantifler™ Human DNA Quantification Kit, AmpF/STR[®] Identifiler™ PCR Amplification Kit, GenerScan™ -500 LIZ[®] Size Standard, AmpF/STR[®] Identifiler™ Allelic Ladder, Optical Adhesive Cover, Plate Septa 96-Well, MicroAmp[®] Caps (8 caps/strip), MicroAmp[®] 96 Well Tray for Tubes with Caps, MicroAmp[®] 96-Well Tray-Retainer Set, MicroAmp[®] Optical 96-Well Reaction Plate, ABI PRISM[®] 7000 Sequence Detection (SDS), GeneAmp[®] PCR System 9700, ABI PRISM[®] 3100 genetic analyzer. Instruments as the Vortex mixer (Labnet VX100), Microcentrifuge (PORVALL[®] *pico*), Heat block (FINEPCR SLBI28) and Autoclave (Astell AMA 240) were used in this study.

2. Volunteers

Ten volunteers (five males and five females) already had DNA profiles for the staff DNA profiles database and worked at the Forensic DNA Service Center, one service center of the Central Institute of Forensic science (CIFS), were asked for helpful in this study. The volunteers were stayed in office and laboratory. Their activities such as did the laboratory, document and computer.

3. Adhesive tape sample preparation

The three trademarks commercial electrical adhesive tape were used as starting material for all experiments: Scotch[®] 3M 1710 Vinyl Electrical Tape, Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape and YAZAKI P.V.C. Tape. These trademarks commercial electrical adhesive tape available in the market, the cost was not expensive and always found at the crime scene. Every time before using, each electrical adhesive tape was cleaned with 80% ethanol and aqua bidest, to remove to foreign DNA, before storing in the sterile clean glass plate.

4. Epithelial cells collection

4.1 Reference sample

The volunteers already had DNA profiles for the staff DNA profiles database. Their epithelial cells were collected by buccal swab method and extracted by Chelex[®] 100 extraction method. The quantity of extracted DNA was quantified by Real-Time PCR, using ABI PRISM[®] 7000 Sequence Detection System (7000 SDS) and Quantifiler[™] Human DNA Quantification kit (Applied Biosystems, CA, USA). After quantitative, the DNA was amplified by using GeneAmp[®] PCR System 9700 and AmpF/STR[®] Identifiler[™] PCR Amplification Kit (Applied Biosystems). The capillary electrophoresis and STRs typing were performed with Data Collection Software Version 2.0 on ABI PRISM[®] 3100 Genetic Analyzer (Applied Biosystems).

4.2 Epithelial cells collection from the volunteers

A total 90 fingerprints from ten different person who working in the office and laboratory were applied to the three trademark electrical adhesive types: Scotch[®] 3M 1710 Vinyl Electrical Tape, Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape and YAZAKI P.V.C. Tape. These electrical adhesive tapes were cut into 10 cm long, then, each volunteer was used their thumbs and index fingers to touch on upper and lower end of the tape. For the orderliness, the fingerprints information were carried out by pressing for 10 sec. Subsequently, the tape was folded over itself and stored in a plastic zip locked bag at room temperature until the process of DNA extraction. Three sets of specimens, each electrical adhesive tape trademark was used for three extraction methods, were collected in the evening without washing the hand and each was used for quantitative and qualitative analysis.

5. DNA extraction

Before DNA extraction, each electrical adhesive tape was removed from the plastic zip locked bag. The tape was cut into small pieces and placed into a sterile 50 ml centrifuge tube. Negative control sample was made with electrical adhesive tape without touching by any person.

5.1 DNA IQ™ System

The extraction procedure was performed following DNA IQ™ System-Database Protocol. For extraction, Four ml of prepared Lysis Buffer were added into a sterile 50 ml centrifuge tube which containing the electrical adhesive tape sample and mixed by the vortex for 10 sec. The sample was incubated in the Lysis Buffer overnight at room temperature. Then, the sample was mixed by vortex 10 sec and incubated at 95 °C for 30 min. The tube was removed from the heat source and transferred the solution to a new labeled sterile 15 ml centrifuge tube. Subsequently, seven µl of DNA IQ resin were added, followed by pulse-vortex and incubated at room temperature for 5 min. Then, the tube was placed in the MagneSphere® Technology Magnetic Separation Stand (Promega, WI, USA). The supernatant was carefully removed and discarded without removing the tube from magnetic stand. 100 µl of prepared Lysis Buffer were added and carefully removed again. After that, 100 µl of 1X Wash Buffer were added to the tube and pipette up and down gently to re-suspend the magnetic beads. The vortex tube was placed in the magnetic stand. The supernatant was carefully removed and discarded. 1X Wash Buffer were added and repeated three times, then, opened the lid of the tube for air-dry at room temperature for 5 min. 15 µl of Elution Buffer were added and incubated at 65 °C for 5 min. Then, the tube was removed from the source, vortexed and placed in magnetic stand. Subsequently, specific volume -15 µl of the supernatant containing the DNA- were removed to a new sterile 1.5 ml microcentrifuge tube and stored at 4 °C in refrigerator.

5.2 Chelex® 100 extraction

The extraction procedure was performed previously described. Four ml of sterile double distilled water were added into a sterile 50 ml centrifuge tube which containing the electrical adhesive tape sample followed by pulse-vortex for 10 sec. The tube was then incubated at room temperature and shook overnight followed by pulse - vortex for 10 sec and centrifuged at 13,000 rpm for 5 min. After that, the supernatant was removed and discarded, but obtained about 30 µl. Subsequently, 200 µl of 5% Chelex stock suspension were added into each tube, followed by 20 µl of 20 mg/ ml of Proteinase K and the tube was pulse-vortex for 10 sec. The tube was incubated

at 56 °C for 30 min, after vortexed at maximum speed for 10 sec the tube was incubated at 100 °C for 8 min. Finally the tube was vortexed at maximum speed for 10 sec and centrifuged at 13,000 rpm for 5 min. 200 µl of supernatant was pipetted into a new sterile 1.5 ml microcentrifuge tube. The Microcon™ YM-100 (Millipore, MA, USA) was used to concentrate samples up to 20 µl and stored at 4 °C in refrigerator.

5.3 Phenol-Chloroform extraction

The extraction procedure was performed using Phenol: Chloroform Kit (PIERCE, CA, USA). Four ml of Extraction buffer (contains 10 mM TRIS-HCl, 100 mM NaCl, 60 mM DTT, 50 mM EDTA, 20% Sodium dodecyl sulfate (20% SDS) and 20 mg/ml Proteinase K) were added into a sterile 50 ml centrifuge tube which containing the electrical adhesive tape sample followed by pulse-vortex for 10 sec. The tube was then incubated overnight in 56 °C water bath followed by pulse-vortex for 10 sec, then, the tube was removed from water bath and spin briefly before opening. Two ml of phenol/chloroform/isoamyl-alcohol were added to the tube and vortexed for 30 sec to achieve a milky emulsion, then, the electrical adhesive tape was removed and subsequently centrifuged at 10,000 rpm for 5 min. Upper aqueous layer were removed into new labeled 2.0 ml microcentrifuge tube, then, an equal volume of phenol/chloroform/isoamyl-alcohol was added to the tube and vortexed for 30 sec. The tube was centrifuged at 10,000 rpm for 5 min and removed upper aqueous layer into new labeled 2.0 ml microcentrifuge tube. The Microcon™ YM-100 (Millipore) was used to concentrate sample up to 20 µl and stored at 4 °C in refrigerator.

6. DNA quantitation

For quantitation the extracted DNA, Quantifiler™ Human DNA Quantification Kit (Applied Biosystems, CA, USA) was used. The kit contained Quantifiler™ PCR Reaction Mix, contained AmpliTag Gold® DNA Polymerase, dNTPs, dUTP, passive reference and buffer components; Quantifiler™ Primer Mix, contained forward and reverse primer, probe and internal PCR control (IPC) system which included primer, probe and template; and Quantifiler™ DNA Standard.

6.1 DNA Standard and sample preparations for Real-Time PCR

The standard curve was developed by preparing four folds dilution series. For dilution, 200 ng/µl Quantifiler™ Human DNA Standard was diluted by sterile TE buffer, containing 10 mM Tris-HCl (ph 8.0) and 0.1 mM Na₂EDTA, to concentration 25, 6.25, 0.5625, 0.3905, 0.0975 and 0.0245 ng/µl.

The master mix containing 10.5 μ l of QuantifilerTM Primer Mix and 12.5 μ l of QuantifilerTM PCR Reaction Mix per sample was pipetted to the tube followed by briefly the vortexed and centrifuged. Subsequently, 23 μ l of the master mix were pipetted into each well of the 96-Well Optical Reaction Plate (Applied Biosystems). After created a plate document – contained sample type information, primer and probe used and sample location on the plate – 2 μ l of each concentration of QuantifilerTM Human DNA Standard was loaded into each well in duplications, then, 2 μ l of the extracted DNA were loaded. Therefore, the total volume in each well was 25 μ l. The bubbles were checked that there are no in each well before the plate was sealed with Optical Adhesive Cover and then a compression pad (Applied Biosystems) was placed over the plate. Finally, this plate was placed in the Real-Time PCR instrument.

6.2 Real-Time PCR

The Real-Time PCR data collection and analysis were performed by using ABI PRISM 7000 Sequence Detection System (7000 SDS) and SDS Software Version 1.0 (Applied Biosystems). The instrument was turned on for 15 min before running and the Real-Time PCR condition were set to three steps that were initial denaturation at 95 °C for 10 min, denaturation at 95 °C for 15 min, annealing and extension at 60 °C for 1 min was repeated 40 times.

6.3 Results recording and interpretation

After analysis Real-Time PCR results, the standard curve's slope, intercept, R^2 and C_T value of the IPC in each well were recorded and referred to PCR efficiency. The human DNA quantity in each sample was recorded in concentration ng/ μ l. After that, assign number "1" to the DNA samples that having the concentration equal or above 0.1 ng/ μ l in order to be referred as "effective sample for PCR", and assign number "0" to the DNA samples that having the concentration less than 0.1 ng/ μ l (Low Copy Number (LCN) DNA concentration) in order to be referred as "ineffective sample for PCR". These numbers were used for statistical analysis.

7. DNA typing by Multiplex Polymerase Chain Reaction

After quantitation, the twelve "effective" extracted DNA samples (given number "1") by methods of DNA IQTM System, Chelex[®] 100 and Phenol: Chloroform Kit of all electrical adhesive tape trademarks were amplified and genotyped. The DNA amplification was performed with AmpF/STR[®] IdentifilerTM PCR Amplification Kit (Applied Biosystems).

7.1 Reaction preparation

A master mix containing 10.0 μl of AmpF/STR[®] PCR Reaction Mix, 5.0 μl of AmpF/STR[®] Identifiler[™] Primer Set and 0.5 μl of Amplitaq Gold[®] DNA Polymerase per sample was performed.

After vortexing, this master mix was aliquoted into PCR tube followed by the DNA template for each sample was added. The “effective” extracted DNA samples (given number “1”) were added to the PCR tubes which containing 15 μl of master mix. The PCR total volume was 25 μl (Table 1).

Table 1 Amounts of template DNA from collected samples in PCR amplification

DNA samples	DNA concentration (ng/ μl)	DNA template Volume (μl)	dH ₂ O (μl)	Total PCR Template (ng)	Reaction Volume (μl)
ReSI105	3.290	0.608	9.392	2.00	25
ReSI109	2.970	0.673	9.327	2.00	25
ReSI209	2.340	0.855	9.145	2.00	25
ReSI306	1.845	1.084	8.916	2.00	25
ReSC205	0.234	8.547	1.453	2.00	25
ReSC209	1.440	1.389	8.611	2.00	25
ReSP101	1.015	1.970	8.030	2.00	25
ReSP105	0.630	3.175	6.825	2.00	25
ReSP204	0.429	4.662	5.338	2.00	25
ReSP207	1.425	1.404	8.596	2.00	25
ReSP305	0.590	3.390	6.610	2.00	25
ReSP307	3.365	0.594	9.406	2.00	25

“ReSI”, “ReSC” and “ReSP” indicated that the adhesive tape extracted by DNA IQ[™] System, Chelex[®] 100 extraction and Phenol-Chloroform extraction method, respectively; the number “1”, “2” and “3” after the alphabet indicated the adhesive tape’s trademark Scotch[®] 3M 1710 Vinyl Electrical Tape, Scotch[®] 3M super 33+ All the weather Vinyl Electrical Tape and YAZAKI P.V.C. Tape, respectively; the next two number indicated to the volunteer number.

7.2 DNA amplification by Multiplex PCR

DNA samples were amplified by using GeneAmp[®] PCR System 9700 (Applied Biosystems). The condition was followed: initial denaturation at 25 °C for 11 min; 28 cycles of denaturation at 94 °C, annealing at 59 °C and extension at 72 °C (one min at each temperature); and the final extension at 60 °C for 60 min. After the final extension step, the temperature was then brought down to 4 °C. The PCR product tubes were covered with foil and stored at -20 °C until genotyping.

8. Capillary electrophoresis and genotyping

8.1 Sample preparation

A master mix containing 0.5 µl of GeneScan[™] -500 LIZ Size Standard (Applied Biosystems) and 11.5 µl of Hi-Di formamide per sample was prepared. After vortexing, 12 µl of master mix were aliquoted into each well of 96-Well Optical Reaction Plate (Applied Biosystems). For each run (Figure 2), 3.5 µl of AmpF/STR[®] Identifiler[™] Allelic Ladder were loaded into the first well, which corresponded to position A1. 1.5 µl of PCR positive (A2) and PCR negative control (B1) were loaded next well. Only the master mix was loaded in well B2 for a blank of electrophoresis. After that, 2 µl of PCR product from each sample was loaded into its well. The plate was sealed with Plate Septa 96-Well (Applied Biosystems) before placed on GeneAmp[®] PCR System 9700 (Applied Biosystems) and heated at 95 °C for 5 min before the temperature was rapidly down to 4 °C and held for 3 min. Then, the plate was placed on plate base and covered with plate retainer.

	1	2	3	4	5	6	7	8	9	10	11	12
A	L	P	L	P	L	P	L	P	L	P	L	P
B	N	R	13	14	27	28	41	42	55	56	69	70
C	1	2	15	16	29	30	43	44	57	58	71	72
D	3	4	17	18	31	32	45	46	59	60	73	74
E	5	6	19	20	33	34	47	48	61	62	75	76
F	7	8	21	22	35	36	49	50	63	64	77	78
G	9	10	23	24	37	38	51	52	65	66	79	80
H	11	12	25	26	39	40	53	54	67	68	81	82

Figure 2 A diagram showing typical positions of allelic ladder markers, PCR positive and negative control, electrophoresis reagent blank and DNA samples on a 96-Well plate.

8.2 Capillary electrophoresis

The capillary electrophoresis and STRs typing was performed with Data Collection Software Version 2.0 on ABI PRISM[®] 3100 Genetic Analyzer. Before operating the instrument, 1X TBE buffer (Applied Biosystems) and distilled water were changed every time.

9. Data analysis

The STRs fragment were analysed with GeneMapper[™] ID Software Version 3.1 on ABI PRISM[®] 3100 Genetic Analyzer (Applied Biosystems). The peak amplitude threshold (PAT) was set at 100 RFU. DNA profiles of AmpF/STR[®] Identifiler[™] Allelic Ladder, size standard and positive control were checked in the process of DNA analysis compared with the reference profile for each run. No peak should be present for negative control analyzed. Then, DNA profiles of the volunteers on electrical adhesive tape were analyzed and matched with reference volunteers DNA profiles.

10. Statistical analysis

The effective and ineffective DNA sample for PCR were analyzed by Two-way (Test of Independence) Chi-Square, separately for DNA extraction method and electrical adhesive tape trademarks. In this study, Test of Independence Chi-Square was used to examine the relationship between subjects' scores on two qualitative or categorical variables. A value of $p < 0.05$ was considered statistically significant. SPSS program for windows was used for all analyses.

CHAPTER IV

RESULTS

1. Different methods of DNA extraction

Extractions of DNA samples from the electrical adhesive tapes lead to conclusion that the DNA could be extracted from the skin epithelial cells left on electrical adhesive tape. Electrical adhesive tapes were collected from 10 volunteers by using 3 trademarks of the commercial electrical adhesive tapes, later on extracted by DNA IQ™ System, Chelex® 100 and Phenol: Chloroform Kit, as previously described in section 3.5. Each scheme of the extracted DNA was quantitated as described in section 3.6.

The quantitation results of extracted DNA lead to the conclusion that different extraction method yields different quantity of DNA. Among three DNA extraction methods; Phenol: Chloroform kit (260.344 ng/μl), and DNA IQ™ System (259.458 ng/μl) were better than Chelex® 100 (54.634 ng/μl). No different DNA yields from Phenol: Chloroform kit and DNA IQ™ System was significantly demonstrated (Table 2 and Figure 3).

Table 2 DNA yield from 10 volunteers after extraction by DNA IQ™ System, Chelex® 100 and Phenol: Chloroform Kit.

DNA Samples	DNA yield (ng/μl)		
	DNA IQ™ System	Chelex® 100	Phenol: Chloroform kit
Volunteer no. 1	0.2594	0.0282	0.4827
Volunteer no. 2	0.2302	0.0210	0.1480
Volunteer no. 3	0.1018	0.0156	0.0917
Volunteer no. 4	0.2373	0.0258	0.3632
Volunteer no. 5	1.4315	0.1562	0.3867
Volunteer no. 6	0.8902	0.0185	1.3490
Volunteer no. 7	0.2987	0.0497	0.5905
Volunteer no. 8	0.1277	0.0723	0.3062
Volunteer no. 9	1.8663	0.4969	0.3917
Volunteer no.10	0.3227	0.0263	0.2295

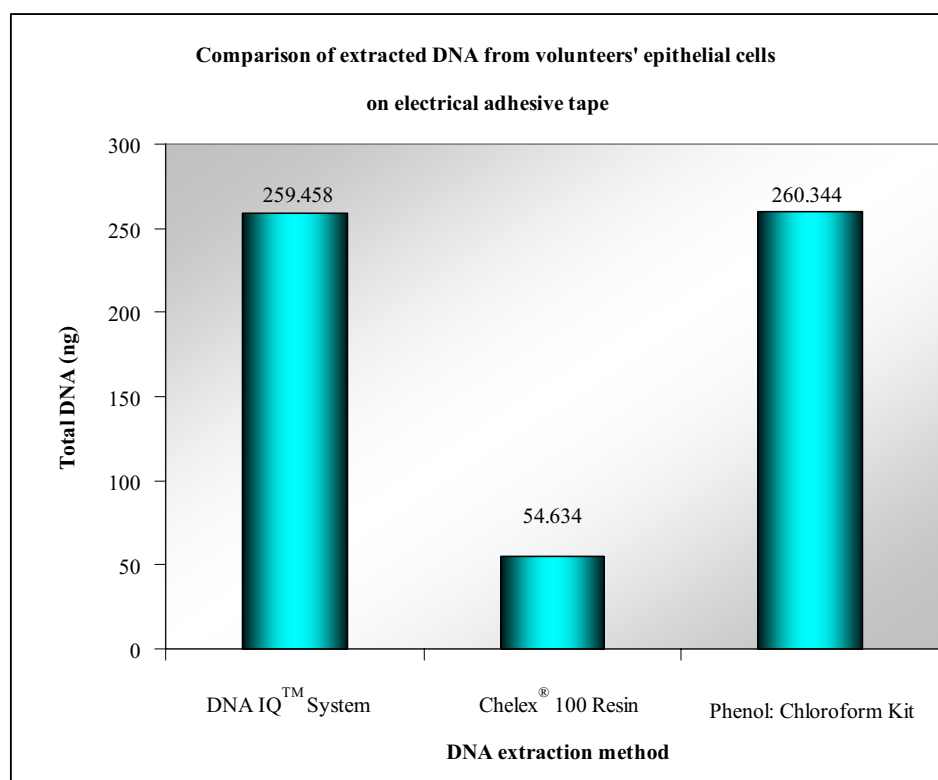


Figure 3 Amounts of DNA on electrical adhesive tape samples from 10 volunteers after extraction by DNA IQ™ System, Chelex® 100 and Phenol: Chloroform Kit.

2. The quantitation of DNA extracted from electrical adhesive tape

All the extracted DNA from each trademark sample of the electrical adhesive tape was quantified as described in section 3.6 previously. After the quantitations by Real-Time PCR method, the results were found as standard curve (Appendix A) and C_T result was shown that R^2 value were 0.999 and 0.997 that were greater than 0.99. The slope of standard curves was between -2.84 and -2.58, which were in the range of -2.9 to 3.3.

Then, measure the concentration of each DNA extraction method, by each adhesive tape trademark sample, in nanogram per microlitre (ng/μl). Among three DNA extraction methods; Phenol: Chloroform kit (260.344 ng/μl), and DNA IQ™ System (259.458 ng/μl) were better than Chelex® 100 (54.634 ng/μl). No different DNA yields from Phenol: Chloroform kit and DNA IQ™ System was significantly demonstrated as shown in Figure 3.

After that, assign number “1” to the DNA samples that having the concentration equal or above 0.1 ng/μl in order to be referred as “effective sample for PCR”, and assign number “0” to

the DNA samples that having the concentration less than 0.1 ng/μl in order to be referred as “ineffective sample for PCR” (Appendix B). The summary results were as following; extracted DNA samples from both DNA IQ™ System and Phenol: Chloroform Kit, were “effective” at 25 out of 30, or 83.33 %. The extracted DNA samples from Chelex® 100 were “effective” at 4 out of 30, or 13.33 %. And, the extracted DNA samples from each electrical adhesive tape trademark were “effective” at 18 out of 30, or 60.00 %.

3. Statistical analysis

All data was analyzed by SPSS program for windows. The statistical test methods are two-ways Chi-square test with p -value less than 0.05. These are to indicate the statistical significance relationship between independent and dependent variables. In this study, there are two hypotheses.

3.1 Hypothesis I

Research Hypothesis: The DNA extraction methods relate to the efficiency in PCR process.

Statistical Hypothesis: $H_0: r = 0$

$H_1: r \neq 0$

Analysis of association between DNA extraction method and PCR efficiency was found that it had statistically significance ($\chi^2 = 40.833$; $df = 2$; p -value = 0.000). These statistically significance indicated that the PCR efficiency depended on DNA extraction methods (Table 3 and Table 4).

Table 3 Statistical test showed the effect of DNA extraction method on PCR efficiency

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	40.833 ^a	2	.000
Likelihood Ratio	43.514	2	.000
Linear-by-Linear Association	.000	1	1.000
N of Valid Cases	90		

a. 0 cells (.0%) have expected count less than 5. The minimum expected count is 12.00.

Table 4 Standardized residual of DNA extraction methods

			Analysis results		
			Have not PCR	Have PCR	Total
			efficiency	efficiency	
DNA extraction method	DNA IQ System	Count	5	25	30
		Expected Count	12.0	18.0	30.0
		Residual	-7.0	7.0	
		Std. Residual	-2.0	1.6	
	Chelex 100	Count	26	4	30
		Expected Count	12.0	18.0	30.0
		Residual	14.0	-14.0	
		Std. Residual	4.0	-3.3	
	Phenol: Chloroform Kit	Count	5	25	30
		Expected Count	12.0	18.0	30.0
		Residual	-7.0	7.0	
		Std. Residual	-2.0	1.6	
	Total	Count	36	54	90
		Expected Count	36.0	54.0	90.0

We found that the Chelex[®] 100 DNA extraction method had less PCR efficiency than expectation (Standardize residual=-3.3). And there was no PCR efficiency more than expectation, too (Standardize residual=4.0). In the other hand, the DNA extraction methods of DNA IQ[™] System and Phenol: Chloroform Kit had PCR efficiency more than expectation (Standardize residual=1.6), and not less than expectation (Standardize residual=-2.0).

3.2 Hypothesis II

Research Hypothesis: The trademarks of electrical adhesive tape relate to the efficiency in PCR process.

Statistical Hypothesis: $H_0: r = 0$

$H_1: r \neq 0$

For the PCR efficiency analysis results of each trademark of the electrical adhesive tapes were shown as no significant differences among all of them (Scotch[®] 3M 1710 Vinyl Electrical Tape, Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape and YAZAKI P.V.C. Tape) and the PCR efficiency ($\chi^2 = 0.000$; $df = 2$; $p\text{-value} = 1.000$). This statistical analysis is shown in Table 5.

Table 5 Statistical test showed the effect of electrical adhesive tape trademarks on PCR efficiency

	Value	df	Asymp. Sig. (2-sided)
Pearson Chi-Square	.000 ^a	2	1.000
Likelihood Ratio	.000	2	1.000
Linear-by-Linear Association	.000	1	1.000
N of Valid Cases	90		

a. 0 cells (.0%) have expected count less than 5. The minimum expected count is 12.00.

4. DNA profiles of collected samples

After measuring the DNA quantity extracted from the Real-Time PCR method, the twelve effective samples of extracted DNA by all 3 methods and all 3 trademarks were amplified and analyzed as previously described in section 3.7 and 3.8, respectively. The electrophoregrams (Appendix D) were analyzed to compare with the DNA profile reference of each volunteer kept in laboratory (Appendix C). DNA profiles of Allelic ladder, size standard, positive control and negative control were checked every time for DNA samples analysis.

The DNA profile reference collected from volunteers' buccal swabs, were showed in Table 8-17. And, the volunteers' DNA profiles extracted from the touched electrical adhesive tape, were

showed in Table 18-29. The full DNA profile described as allele presents in every locus of 15 STR loci. The homologous allele shows one allele on each locus while the heterozygous allele shows two alleles at a locus.

Representative electrophoregram was showed in Figure 4, an alleles was presented at all loci as full DNA profile. The blue, green, black and red peak indicated that labeled the amplified DNA product with 6-FAM™, VIC™, NED™ and PET™, respectively. The homologous alleles were examined at D19S433, TPOX and AMEL loci while the heterozygous alleles were showed at D8S1179, D6S11, D7S820, CSF1PO, D3S1358, TH01, D13S539, D2S1338, vWA, D18S51, D5S818 and FGA loci.

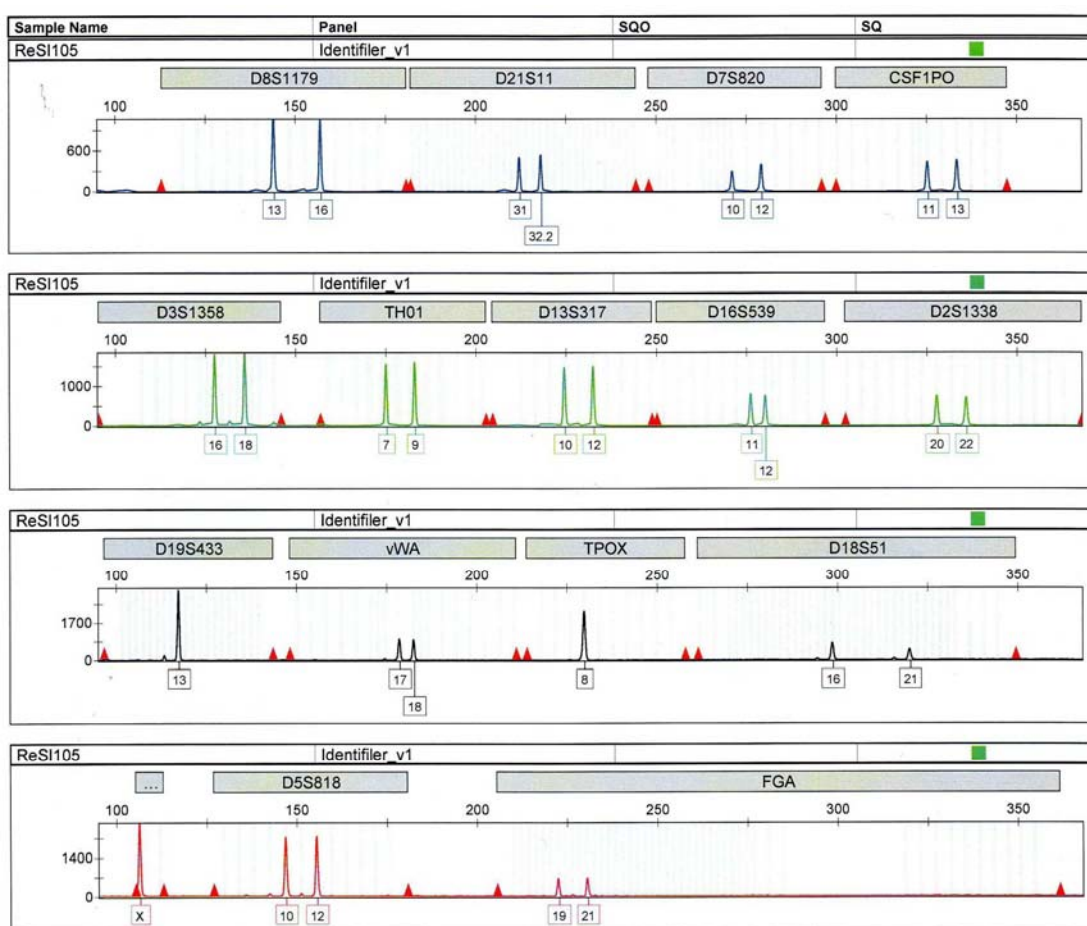


Figure 4 Representative electrophoregram from volunteer no.5.

The full STR profile of the volunteers no.1, 4, 5, 6, 7 and 9 (Table 18-29) showed that alleles were obtained at all loci and matches with the DNA profiles derived from buccal swabs as the reference sample of each volunteer (Table 6 and 7).

Table 6 Comparative reference DNA profiles and sample DNA profiles of volunteer no. 1, 4 and 5.

Marker	Reference DNA Profile of volunteer no.1	DNA Profile from sample of volunteer no.1	Reference DNA Profile of volunteer no.4	DNA Profile from sample of volunteer no.4	Reference DNA Profile of volunteer no.5	DNA Profile from sample of volunteer no.5
D8S1179	11,13	11,13	10,10	10,10	13,16	13,16
D21S11	32.2,32.2	32.2,32.2	28,29	28,29	31,32.2	31,32.2
D7S820	11,12	11,12	11,11	11,11	10,12	10,12
CSF1PO	11,11	11,11	12,12	12,12	11,13	11,13
D3S1358	16,16	16,16	15,16	15,16	16,18	16,18
TH01	6,10	6,10	7,7	7,7	7,9	7,9
D13S317	8,11	8,11	8,13	8,13	10,12	10,12
D16S539	9,11	9,11	11,12	11,12	11,12	11,12
D2S1338	17,24	17,24	24,24	24,24	20,22	20,22
D19S433	14,14	14,14	13,14.2	13,14.2	13,13	13,13
vWA	14,16	14,16	14,17	14,17	17,18	17,18
TPOX	8,9	8,9	8,8	8,8	8,8	8,8
D18S51	13,14	13,14	12,15	12,15	16,21	16,21
AMEL	X,X	X,X	X,X	X,X	X,X	X,X
D5S818	9,13	9,13	10,11	10,11	10,12	10,12
FGA	22,25	22,25	22,23	22,23	19,21	19,21

Table 7 Comparative reference DNA profiles and sample DNA profiles of volunteer no. 6, 7 and 9.

Marker	Reference DNA Profile of volunteer no.6	DNA Profile from sample of volunteer no.6	Reference DNA Profile of volunteer no.7	DNA Profile from sample of volunteer no.7	Reference DNA Profile of volunteer no.9	DNA Profile from sample of volunteer no.9
D8S1179	16,17	16,17	11,15	11,15	13,14	13,14
D21S11	29,32.2	29,32.2	30,34.2	30,34.2	29,29	29,29
D7S820	8,10	8,10	8,11	8,11	10,11	10,11
CSF1PO	12,13	12,13	12,12	12,12	10,14	10,14
D3S1358	15,15	15,15	17,18	17,18	16,18	16,18
TH01	6,9,3	6,9,3	8,9,3	8,9,3	6,10	6,10
D13S317	13,14	13,14	8,11	8,11	8,8	8,8
D16S539	9,11	9,11	11,12	11,12	11,11	11,11
D2S1338	18,18	18,18	17,24	17,24	18,23	18,23
D19S433	13,14.2	13,14.2	13,13.2	13,13.2	14.2,16.2	14.2,16.2
vWA	17,18	17,18	14,19	14,19	17,17	17,17
TPOX	8,9	8,9	8,8	8,8	8,8	8,8
D18S51	13,15	13,15	13,18	13,18	12,14	12,14
AMEL	X,Y	X,Y	X,Y	X,Y	X,Y	X,Y
D5S818	12,13	12,13	10,10	10,10	13,13	13,13
FGA	25,25	25,25	22,22	22,22	21,22	21,22

Table 8 STR profile of the DNA from buccal swab for reference sample of volunteer no.1

Sample Name: -

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	11	13
D21S11	32.2	32.2
D7S820	11	12
CSF1PO	11	11
D3S1358	16	16
TH01	6	10
D13S317	8	11
D16S539	9	11
D2S1338	17	24
D19S433	14	14
vWA	14	16
TPOX	8	9
D18S51	13	14
AMEL	X	X
D5S818	9	13
FGA	22	25

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 9 STR profile of the DNA from buccal swab for reference sample of volunteer no.2

Sample Name: -

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	15	15
D21S11	29	30
D7S820	11	12
CSF1PO	10	11
D3S1358	15	16
TH01	7	9
D13S317	8	11
D16S539	13	13
D2S1338	19	24
D19S433	13	14
vWA	16	16
TPOX	8	11
D18S51	15	15
AMEL	X	X
D5S818	10	11
FGA	23	23

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 10 STR profile of the DNA from buccal swab for reference sample of volunteer no.3

Sample Name: -

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	12	13
D21S11	29	30
D7S820	8	11
CSF1PO	12	13
D3S1358	16	16
TH01	6	10
D13S317	9	11
D16S539	12	12
D2S1338	23	24
D19S433	14	15
vWA	16	18
TPOX	8	9
D18S51	13	13
AMEL	X	X
D5S818	12	14
FGA	23	23

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 11 STR profile of the DNA from buccal swab for reference sample of volunteer no.4

Sample Name: -

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	10	10
D21S11	28	29
D7S820	11	11
CSF1PO	12	12
D3S1358	15	16
TH01	7	7
D13S317	8	13
D16S539	11	12
D2S1338	24	24
D19S433	13	14.2
vWA	14	17
TPOX	8	8
D18S51	12	15
AMEL	X	X
D5S818	10	11
FGA	22	23

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 12 STR profile of the DNA from buccal swab for reference sample of volunteer no.5

Sample Name: -

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	13	16
D21S11	31	32.2
D7S820	10	12
CSF1PO	11	13
D3S1358	16	18
TH01	7	9
D13S317	10	12
D16S539	11	12
D2S1338	20	22
D19S433	13	13
vWA	17	18
TPOX	8	8
D18S51	16	21
AMEL	X	X
D5S818	10	12
FGA	19	21

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 13 STR profile of the DNA from buccal swab for reference sample of volunteer no.6

Sample Name: -

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	16	17
D21S11	29	32.2
D7S820	8	10
CSF1PO	12	13
D3S1358	15	15
TH01	6	9.3
D13S317	13	14
D16S539	9	11
D2S1338	18	18
D19S433	13	14.2
vWA	17	18
TPOX	8	9
D18S51	13	15
AMEL	X	Y
D5S818	12	13
FGA	25	25

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 14 STR profile of the DNA from buccal swab for reference sample of volunteer no.7

Sample Name: -

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	11	15
D21S11	30	34.2
D7S820	8	11
CSF1PO	12	12
D3S1358	17	18
TH01	8	9.3
D13S317	8	11
D16S539	11	12
D2S1338	17	24
D19S433	13	13.2
vWA	14	19
TPOX	8	8
D18S51	13	18
AMEL	X	Y
D5S818	10	10
FGA	22	22

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 15 STR profile of the DNA from buccal swab for reference sample of volunteer no.8

Sample Name: -

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	10	16
D21S11	31.2	33.2
D7S820	8	12
CSF1PO	11	12
D3S1358	17	18
TH01	9	9
D13S317	8	13
D16S539	12	12
D2S1338	18	19
D19S433	14	16
vWA	17	17
TPOX	8	11
D18S51	13	18
AMEL	X	Y
D5S818	13	15
FGA	22	23.2

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 16 STR profile of the DNA from buccal swab for reference sample of volunteer no.9

Sample Name: -

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	13	14
D21S11	29	29
D7S820	10	11
CSF1PO	10	14
D3S1358	16	18
TH01	6	10
D13S317	8	8
D16S539	11	11
D2S1338	18	23
D19S433	14.2	16.2
vWA	17	17
TPOX	8	8
D18S51	12	14
AMEL	X	Y
D5S818	13	13
FGA	21	22

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 17 STR profile of the DNA from buccal swab for reference sample of volunteer no.10

Sample Name: -

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	12	15
D21S11	30	32.2
D7S820	12	12
CSF1PO	11	12
D3S1358	15	16
TH01	8	8
D13S317	8	8
D16S539	10	12
D2S1338	20	24
D19S433	12	13
vWA	17	19
TPOX	8	11
D18S51	13	14
AMEL	X	Y
D5S818	11	12
FGA	22	22.2

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center,
CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center,
 2008), n. pag.

Table 18 STR profile of the DNA from DNA IQ™ System extraction (ReSI) collected by Scotch® 3M 1710 Vinyl Electrical Tape of volunteer no.5

Sample Name: ReSI105

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	13	16
D21S11	31	32.2
D7S820	10	12
CSF1PO	11	13
D3S1358	16	18
TH01	7	9
D13S317	10	12
D16S539	11	12
D2S1338	20	22
D19S433	13	13
vWA	17	18
TPOX	8	8
D18S51	16	21
AMEL	X	X
D5S818	10	12
FGA	19	21

Table 19 STR profile of the DNA from DNA IQ™ System extraction (ReSI) collected by Scotch® 3M 1710 Vinyl Electrical Tape of volunteer no.9

Sample Name: ReSI109

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	13	14
D21S11	29	29
D7S820	10	11
CSF1PO	10	14
D3S1358	16	18
TH01	6	10
D13S317	8	8
D16S539	11	11
D2S1338	18	23
D19S433	14.2	16.2
vWA	17	17
TPOX	8	8
D18S51	12	14
AMEL	X	Y
D5S818	13	13
FGA	21	22

Table 20 STR profile of the DNA from DNA IQ™ System extraction (ReSI) collected by Scotch® 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.9

Sample Name: ReSI209

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	13	14
D21S11	29	29
D7S820	10	11
CSF1PO	10	14
D3S1358	16	18
TH01	6	10
D13S317	8	8
D16S539	11	11
D2S1338	18	23
D19S433	14.2	16.2
vWA	17	17
TPOX	8	8
D18S51	12	14
AMEL	X	Y
D5S818	13	13
FGA	21	22

Table 21 STR profile of the DNA from DNA IQ™ System extraction (ReSI) collected by YAZAKI P.V.C. Tape of volunteer no.6

Sample Name: ReSI306

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	16	17
D21S11	29	32.2
D7S820	8	10
CSF1PO	12	13
D3S1358	15	15
TH01	6	9.3
D13S317	13	14
D16S539	9	11
D2S1338	18	18
D19S433	13	14.2
vWA	17	18
TPOX	8	9
D18S51	13	15
AMEL	X	Y
D5S818	12	13
FGA	25	25

Table 22 STR profile of the DNA from Chelex[®] 100 extraction (ReSC) collected by Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.5

Sample Name: ReSC205

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	13	16
D21S11	31	32.2
D7S820	10	12
CSF1PO	11	13
D3S1358	16	18
TH01	7	9
D13S317	10	12
D16S539	11	12
D2S1338	20	22
D19S433	13	13
vWA	17	18
TPOX	8	8
D18S51	16	21
AMEL	X	X
D5S818	10	12
FGA	19	21

Table 23 STR profile of the DNA from Chelex[®] 100 extraction (ReSC) collected by Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.9

Sample Name: ReSC209

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	13	14
D21S11	29	29
D7S820	10	11
CSF1PO	10	14
D3S1358	16	18
TH01	6	10
D13S317	8	8
D16S539	11	11
D2S1338	18	23
D19S433	14.2	16.2
vWA	17	17
TPOX	8	8
D18S51	12	14
AMEL	X	Y
D5S818	13	13
FGA	21	22

Table 24 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by Scotch[®] 3M 1710 Vinyl Electrical Tape of volunteer no.1

Sample Name: ReSP101

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	11	13
D21S11	32.2	32.2
D7S820	11	12
CSF1PO	11	11
D3S1358	16	16
TH01	6	10
D13S317	8	11
D16S539	9	11
D2S1338	17	24
D19S433	14	14
vWA	14	16
TPOX	8	9
D18S51	13	14
AMEL	X	X
D5S818	9	13
FGA	22	25

Table 25 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by Scotch[®] 3M 1710 Vinyl Electrical Tape of volunteer no.5

Sample Name: ReSP105

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	13	16
D21S11	31	32.2
D7S820	10	12
CSF1PO	11	13
D3S1358	16	18
TH01	7	9
D13S317	10	12
D16S539	11	12
D2S1338	20	22
D19S433	13	13
vWA	17	18
TPOX	8	8
D18S51	16	21
AMEL	X	X
D5S818	10	12
FGA	19	21

Table 26 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.4

Sample Name: ReSP204

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	10	10
D21S11	28	29
D7S820	11	11
CSF1PO	12	12
D3S1358	15	16
TH01	7	7
D13S317	8	13
D16S539	11	12
D2S1338	24	24
D19S433	13	14.2
vWA	14	17
TPOX	8	8
D18S51	12	15
AMEL	X	X
D5S818	10	11
FGA	22	23

Table 27 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.7

Sample Name: ReSP207

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	11	15
D21S11	30	34.2
D7S820	8	11
CSF1PO	12	12
D3S1358	17	18
TH01	8	9.3
D13S317	8	11
D16S539	11	12
D2S1338	17	24
D19S433	13	13.2
vWA	14	19
TPOX	8	8
D18S51	13	18
AMEL	X	Y
D5S818	10	10
FGA	22	22

Table 28 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by YAZAKI P.V.C. Tape of volunteer no.5

Sample Name: ReSP305

Gender: ☐ Male ☒ Female

Marker	Allele 1	Allele 2
D8S1179	13	16
D21S11	31	32.2
D7S820	10	12
CSF1PO	11	13
D3S1358	16	18
TH01	7	9
D13S317	10	12
D16S539	11	12
D2S1338	20	22
D19S433	13	13
vWA	17	18
TPOX	8	8
D18S51	16	21
AMEL	X	X
D5S818	10	12
FGA	19	21

Table 29 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by YAZAKI P.V.C. Tape of volunteer no.7

Sample Name: ReSP307

Gender: ☒ Male ☐ Female

Marker	Allele 1	Allele 2
D8S1179	11	15
D21S11	30	34.2
D7S820	8	11
CSF1PO	12	12
D3S1358	17	18
TH01	8	9.3
D13S317	8	11
D16S539	11	12
D2S1338	17	24
D19S433	13	13.2
vWA	14	19
TPOX	8	8
D18S51	13	18
AMEL	X	Y
D5S818	10	10
FGA	22	22

CHAPTER V

CONCLUSIONS, DISCUSSIONS AND RECOMMENDATIONS

5.1 CONCLUSIONS

In this study, there are three DNA extraction methods to be assessed and to be determined which one is best suited to recover DNA from electrical adhesive tape, DNA IQ™ System, Chelex® 100 and Phenol: Chloroform kit. DNA recovery from commercial three electrical adhesive tape trademarks, Scotch® 3M 1710 Vinyl Electrical Tape, Scotch® 3M Super 33+ All Weather Vinyl Electrical Tape and YAZAKI P.V.C. Tape, were compared in the same ways. Twelve extracted DNA samples were chosen to create the profile and to compare the quality as well as quantity of the DNA extracted.

The main objectives of this study were to determine the relationships among DNA extraction methods and its efficiency in PCR processes, determine the relationships among various trademarks of electrical adhesive tape and its efficiency in PCR processes, and study problems and to develop recommendations, for the DNA extraction from electrical adhesive tape evidence.

Electrical adhesive tapes were collected from 10 volunteers by using 3 trademarks of the commercial electrical adhesive tapes. These electrical adhesive tapes were cut into 10 cm long, then, each volunteer was used their thumbs and index fingers to touch on upper and lower end of the tape. The fingerprints information were carried out by pressing for 10 sec. Subsequently, the tape was folded over itself and stored in a plastic zip locked bag, later on extracted by DNA IQ™ System, Chelex® 100 and Phenol: Chloroform Kit.

After DNA extraction, the extracted DNA samples were quantified by Real-Time PCR by measuring concentration in ng/μl. All the three DNA extraction methods had been evaluated in this study to identify the quantity difference of all electrical adhesive tape. Among three DNA extraction methods; Phenol: Chloroform kit (260.344 ng/μl), and DNA IQ™ System (259.458 ng/μl) were better than Chelex® 100 (54.634 ng/μl). No different DNA yields from Phenol: Chloroform kit and DNA IQ™ System was significantly demonstrated.

After that, the DNA samples with concentration equal to 0.1 ng/μl or above, were given the number “1”, in order to be referred as “effective” PCR. And, the DNA samples with concentration lower than 0.1 ng/μl -defined as low copy number (LCN) DNA samples- were given the number “0”, in order to be referred as “ineffective” PCR (Appendix B).

Here is the summary result. The extracted DNA from both DNA IQ™ System and Phenol: Chloroform Kit were recovered with PCR efficiency at the rate of 83.33 %. The extracted DNA from Chelex® 100 was recovered with PCR efficiency at the rate of 13.33 %. The extracted DNA from each trademark of the electrical adhesive tapes were recovered with PCR efficiency at the rate of 60.00 %. And all the extracted DNA from this study were recovered with PCR efficiency at 54 out of 90 samples, or equal to 60.00 %.

This study used Chi-Square test as a tool to test the association between the dependent and independent variables. The criteria of statistical significance was set at the *p*-value for lower than 0.05. There was a statistically significant association between DNA extraction methods and PCR efficiency ($\chi^2 = 40.833$; *p*-value = 0.000). It was concluded that DNA extraction method affected the PCR efficiency. Extraction by DNA IQ™ System and Phenol: Chloroform Kit were observed to have PCR efficiency more than expected (Standardized residual = 1.6). For Chelex® 100 extraction, the extracted DNA ad PCR efficiency less than expected (Standardized residual = -3.3).

The statistical analysis results by trademarks of the electrical adhesive tape, showed no statistical significance of the association between different trademarks of the electrical adhesive tape and the PCR efficiency on extraction of DNA samples.

Other than the quantitative analysis results, the twelve “effective” extracted DNA samples (given number “1”) by methods of DNA IQ™ System, Chelex® 100 and Phenol: Chloroform Kit of all electrical adhesive tape trademarks were amplified and genotyped. The electrophoregrams were generated after amplified PCR fragments of DNA samples were separated and analyzed for matching with their reference DNA profiles of each volunteer kept in the laboratory. The full DNA profile of each sample was presented and matched with their reference DNA profiles. There is no mixed DNA profile in this study.

5.2 DISCUSSIONS

Adhesive tape is always recovered at crime scenes, especially in the explosion scenes. Almost every scene, all kinds of adhesive tape may have been used to bind a victim or to adjoin two objects together. Characteristic of the both end of tape can identify some facts related to the case such as the smooth and neat edge cuts may tell that it had been cut by scissor or knife, while the rough and pulling force may tell us that the tape had been pulled until it is broke.

In the explosion scenes, electrical adhesive tape were used to adjoin between explosive and electrical circuit in explosive assembly, caused the explosive workable. Usually, the electrical adhesive tape is torn by grasping it between the index fingers and the thumbs of both hands with thumbs touching. Pulling the hands apart stretches the tape until it was broke, cause the epithelial cells on latent fingerprints of the user always leavings on the electrical adhesive tape.

The explosion creates three kinds of effect: blast pressure, fragmentation, and thermal or heat effects. Blast pressure is the most serious damaged, but thermal or temperature effects are generally the least damaging effects. Heat effect can destroy the DNA a little bit, or cannot do that at all. So, user's epithelial cells were found on electrical adhesive tape and applicable for DNA extraction to identify the person.

In order to create the DNA profiles for person identification, each testing procedure including sample recovery, extraction, amplification and separation. Every step is all-important. One of the most important procedure is DNA extraction. The DNA extraction methods have been developed to separate proteins and other cellular materials from the DNA molecules. Nowadays, there are many methods of DNA extraction and various procedures for get the most DNA quantity from the evidence without contamination that cause unsuccessful DNA typing.

In this study, there are three DNA extraction methods to be assessed and to be determined which one is best suited to recover DNA from electrical adhesive tape, DNA IQ™ System, Chelex® 100 and Phenol: Chloroform kit. DNA recovery from commercial three electrical adhesive tape trademarks, Scotch® 3M 1710 Vinyl Electrical Tape, Scotch® 3M Super 33+ All Weather Vinyl Electrical Tape and YAZAKI P.V.C. Tape, were compared in the same ways.

In this experiment, extracted DNA by DNA IQ™ System gave high amount of DNA (259.458 ng). The DNA IQ™ System uses high efficiency lysis buffer to lyses cells and paramagnetic resin to capture a consistent amount of DNA. The solution impurities can easily be removed by drawing the liquid off the beads and washed multiple times to more thoroughly clean

the DNA. These extraction methods take less time, easy, and high yield of extracted DNA. For all extractions, the electrical adhesive tape was cut into small pieces and extraction was carried out on the whole electrical adhesive tape, caused to use more lysis buffer in the cell lysis process. Therefore, DNA IQ™ System cost was so expensive than other extraction methods.

Extracted DNA by Phenol: Chloroform Kit was yielded high amount of DNA (260.344 ng), due to an adhesive compositions of the electrical adhesive tape called butyl rubber copolymer. When the Phenol: Chloroform was added into the sample tubes, butyl rubber copolymer was melted by chloroform and cause the user's epithelial cells get free from electrical adhesive tape more than other DNA extraction methods. Phenol: chloroform mixture was added to separate the proteins from DNA. The organic extraction method works well for recovery of high molecular weight DNA, removed more contaminants and yielded highest purity of extracted DNA. But, it is time-consuming and involves with hazardous chemicals. The extracted DNA with Phenol: Chloroform Kit had been concentrated by Microcon™ YM-100 (Appendix E) at last.

In this study, the extracted DNA by Chelex® 100 was yielded the lowest amount of DNA (54.634 ng). The electrical adhesive tape samples were incubated in the sterile double distilled water, so epithelial cells could not get free from the electrical adhesive tape. The solution impurities were not removed by this method. The major advantages of this method are quick, simple, very low cost, and not involved with hazardous chemicals. The extracted DNA by Chelex® 100 has been concentrated with Microcon™ YM-100, but the concentration did not yield good results. Chelex® 100 yields rather crude extracted DNA, the remainder impurities might possible to blocked the pores on Microcon™ YM-100.

The statistical analysis results by trademarks of the electrical adhesive tape, showed no statistical significance of the association between different trademarks of the electrical adhesive tape and the PCR efficiency on extraction of DNA samples. Comparison of the sticky on each electrical adhesive tape trademark, from high to low, were Scotch® 3M 1710 Vinyl Electrical Tape, Scotch® 3M Super 33+ All Weather Vinyl Electrical Tape and YAZAKI P.V.C. Tape, respectively. Difference in sticky properties of each trademark might yield different amount of epithelial cells left on the tape, more sticky electrical adhesive tape might yielded more epithelial cells, but it also cannot get free from the tape, because it has more sticky properties. On the contrary, electrical adhesive tape that has a little sticky properties might yielded less epithelial cells, but can get free from the tape because it has a little more sticky properties. So, the statistical

analysis results concluded that there was no statistical significance in association with the trademarks of different electrical adhesive tapes.

Other than the quantitative analysis results, the twelve “effective” extracted DNA samples were amplified and genotyped. The electrophoregrams were generated after amplified PCR fragments of DNA samples were separated and analyzed for matching with their reference DNA profiles of each volunteer kept in the laboratory. The full DNA profile of each sample was presented and matched with their reference DNA profiles. Mixed DNA profile may occur from secondary transfer, the DNA may be transferred from object back to hand. There is no mixed DNA profile in this study.

In order to identify the samples recovered from crime scene, the PCR inhibitor is one of an important concern. DNA extraction is an importance step that can remove contaminants and various ions from explosions. The experimental results obtained in this study show that it is possible to type DNA from electrical adhesive tape.

In this study, the adhesive components could not be the reason for PCR inhibition. This finding is consistent with prior research by Schulz and Reichert (2002), which tried to assess the potential use of latent fingerprints as a DNA source for STR typing and proved possible to type fingerprints removed from the surface with scotch tape and showed that magnetic powder, soot powder and scotch tape not disturbed DNA amplification.

It is also consistent with a study conducted by Li and Harris (2003), which used hydrophilic adhesive tape for collection of evidence for forensic DNA analysis and recovered that DNA profiles can be obtained using hydrophilic adhesive tape without an inhibitory effect on DNA amplification. In addition, Lempan (2007) tried to investigated DNA recovery from forensic clothing samples by tape-lift. These results demonstrated that DNA typing from tape-lift was possible and adhesive components could not inhibitory effect on DNA amplification.

From the results, it can be applied to the forensic cases as; the most suitable method to extract DNA from electrical adhesive tape evidence are using DNA IQ™ System and Phenol: Chloroform extraction method. DNA IQ™ System is the method given high amount of DNA, but more expensive than other DNA extraction method. While, Phenol: Chloroform extraction method is the method given high amount and best quality of DNA. But it involves of using the hazardous chemicals. Therefore, finding of the uses of other chemicals to melt the adhesive compositions of electrical adhesive tape, is interesting and suggested for the further studies.

In this study, the majority of DNA lost could be caused by epithelial cells were not get free from electrical adhesive tape. In the crime scenes, it is possible to discover both spread out and folded electrical adhesive tape. The collecting of biological evidence is so importance to DNA typing. In this study, the electrical adhesive tapes were folded over itself, and then stored in a plastic zip locked bag at room temperature until the time to process for DNA extraction. When spread the tape out, the epithelial cells may be packed or being covered by adhesive or glue components and cannot get free from electrical adhesive tape. Therefore, the other collecting biological evidence method is suggested. The spread electrical adhesive tape should be collected in an evidence box in order to avoid packing of the electrical adhesive tape and to get more epithelial cells free from the adhesive of the electrical adhesive tapes. This could be used to generate further hypotheses in the further studies to understand and to evaluate of the collecting methods for electrical adhesive tape evidence, too.

This study demonstrate the DNA extraction method using the Phenol: Chloroform and DNA IQ™ System as the effective DNA extraction method of the epithelial cells from electrical adhesive tape. At the crime scenes in the three borderland provinces in the South of Thailand, these electrical adhesive tapes are very important forensic evidences that can be linked to the suspect. These will benefit the further justice processes.

5.3 RECOMMENDATIONS

5.3.1 Phenol: Chloroform extraction method involves of using the hazardous chemicals. Therefore, finding of the uses of other chemicals to melt the adhesive compositions of electrical adhesive tape, is interesting and suggested for the further studies.

5.3.2 In this study, the electrical adhesive tapes were folded over itself until the time to process for DNA extraction. When spread the tape out, the epithelial cells may be packed or being covered by adhesive or glue components. Therefore, the spread electrical adhesive tape should be collected in an evidence box in order to avoid packing of the electrical adhesive tape. This could be used to generate further hypotheses in the further studies to understand and to evaluate of the collecting methods for electrical adhesive tape evidence.

5.3.3 For all extractions, the electrical adhesive tape was cut into small pieces and extraction was carried out on the whole electrical adhesive tape, caused to use more lysis buffer in the cell lysis process. For this reason, DNA IQ™ System cost was so expensive than other extraction

methods. Therefore, finding of the uses of other sampling techniques is interesting and suggested. The electrical adhesive tape can be use swabbing techniques in order to avoid using more lysis buffer in the cell lysis process. This could be used to generate further hypotheses in the further studies to understand and to evaluate of the sampling techniques for electrical adhesive tape evidence.

5.3.4 In this study, electrical adhesive tapes were collected by using 3 trademarks of the commercial electrical adhesive tapes. Therefore, finding of the uses of other electrical adhesive tapes trademarks is interesting and suggested for the further studies.

5.3.5 In this study, electrical adhesive tapes were collected from 10 volunteers. Therefore, using more numbers of the volunteer is interesting and suggested for the further studies.

5.3.6 In this study, all electrical adhesive tapes were black. For all DNA extractions, the color of electrical adhesive tape was not extracted. Therefore, finding of the uses of other colors of electrical adhesive tape is interesting and suggested. Difference in colors of each trademark might difference in PCR inhibition. This could be used to generate further hypotheses in the further studies to understand and to evaluate of the colors for electrical adhesive tape evidence.

5.3.7 For the most benefit, latent fingerprints on electrical adhesive tape can be detect before the DNA typing. Because, the DNA typing is possible after treatment with classical latent fingerprint techniques. It is interesting and suggested for the further studies.

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APPENDIX

APPENDIX A

STANDARD CURVE OF REAL-TIME PCR RESULT FROM EXTRACTED DNA OF THE ELECTRICAL ADHESIVE TAPE QUANTITATION

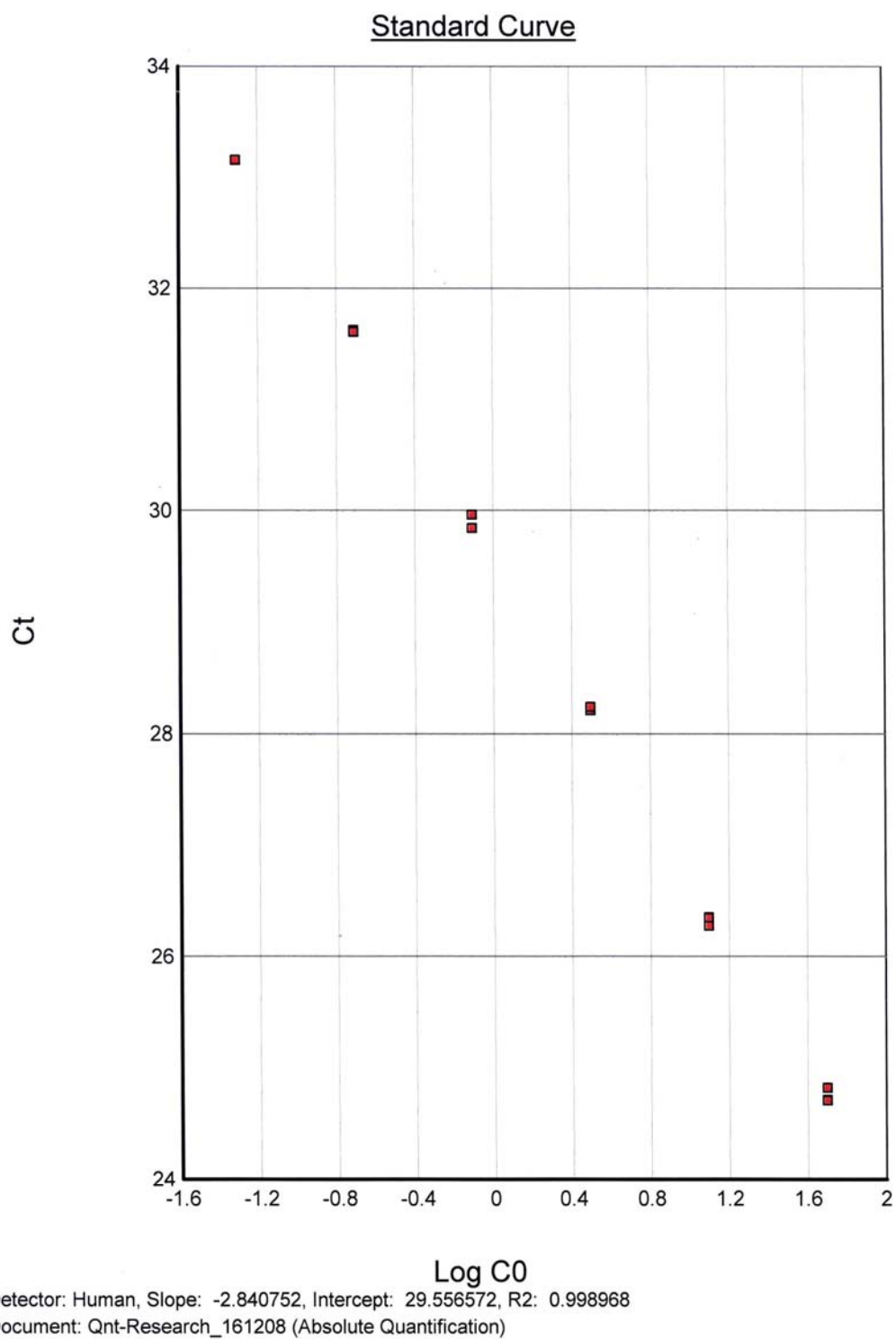


Figure 5 Standard curve of Real-Time PCR result from extracted DNA of the electrical adhesive tape quantitation, slope is -2.84, intercept is 29.557 and R^2 value is 0.999

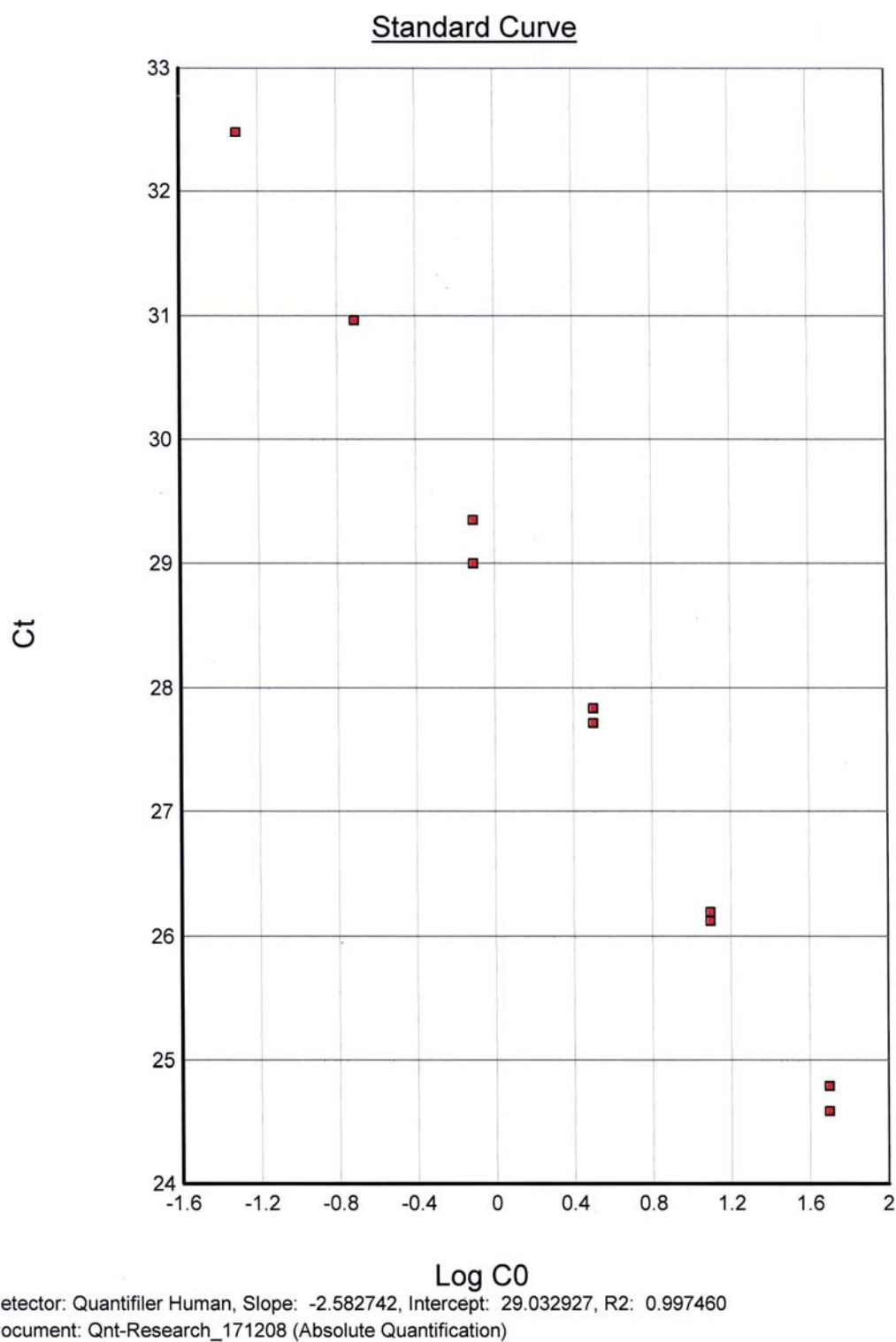


Figure 6 Standard curve of Real-Time PCR result from extracted DNA of the electrical adhesive tape quantitation, slope is -2.58, intercept is 29.033 and R^2 value is 0.997

APPENDIX B

CONCENTRATION OF EXTRACTED DNA FROM ELECTRICAL ADHESIVE TAPE

Table 30 Concentration of extracted DNA from electrical adhesive tape

DNA samples	DNA concentration (ng/μl)	PCR efficiency ("0"= Ineffective, "1"= Effective)
ReSI101	0.1080	1
ReSI102	0.1135	1
ReSI103	0.0815	0
ReSI104	0.2215	1
ReSI105	3.2900	1
ReSI106	0.2105	1
ReSI107	0.3670	1
ReSI108	0.1895	1
ReSI109	2.9700	1
ReSI110	0.5400	1
ReSI201	0.0002	0
ReSI202	0.5150	1
ReSI203	0.0660	0
ReSI204	0.3235	1
ReSI205	0.7200	1
ReSI206	0.6150	1
ReSI207	0.1860	1
ReSI208	0.1260	1
ReSI209	2.3400	1
ReSI210	0.1220	1

“ReSI” indicated that the adhesive tape extracted by DNA IQ™ System, the number “1” and “2” after the alphabet indicated the adhesive tape’s trademark Scotch® 3M 1710 Vinyl Electrical Tape and Scotch® 3M super 33+ All the weather Vinyl Electrical Tape, and the next two number indicated that the volunteer number.

Table 30 Concentration of extracted DNA from electrical adhesive tape (continued)

DNA samples	DNA concentration (ng/μl)	PCR efficiency ("0"= Ineffective, "1"= Effective)
ReSI301	0.6700	1
ReSI302	0.0620	0
ReSI303	0.1580	1
ReSI304	0.1670	1
ReSI305	0.2845	1
ReSI306	1.8450	1
ReSI307	0.3430	1
ReSI308	0.0675	0
ReSI309	0.2890	1
ReSI310	0.3060	1
ReSC101	0.0123	0
ReSC102	0.0014	0
ReSC103	0.0046	0
ReSC104	0.0179	0
ReSC105	0.0630	0
ReSC106	0.0394	0
ReSC107	0.0830	0
ReSC108	0.0322	0
ReSC109	0.0206	0
ReSC110	0.0216	0

“ReSI” and “ReSC” indicated that the adhesive tape extracted by DNA IQ™ System and Chelex® 100 extraction, respectively; the number “1” and “3” after the alphabet indicated the adhesive tape’s trademark Scotch® 3M 1710 Vinyl Electrical Tape and YAZAKI P.V.C. Tape, respectively; the next two number indicated that the volunteer number.

Table 30 Concentration of extracted DNA from electrical adhesive tape (continued)

DNA samples	DNA concentration (ng/μl)	PCR efficiency ("0"= Ineffective, "1"= Effective)
ReSC201	0.0448	0
ReSC202	0.0310	0
ReSC203	0.0328	0
ReSC204	0.0397	0
ReSC205	0.2340	1
ReSC206	0.0160	0
ReSC207	0.0313	0
ReSC208	0.0393	0
ReSC209	1.4400	1
ReSC210	0.0058	0
ReSC301	0.0274	0
ReSC302	0.0306	0
ReSC303	0.0094	0
ReSC304	0.0199	0
ReSC305	0.1715	1
ReSC306	0.0002	0
ReSC307	0.0349	0
ReSC308	0.1455	1
ReSC309	0.0301	0
ReSC310	0.0515	0

“ReSC” indicated that the adhesive tape extracted by Chelex[®] 100 extraction, the number “2” and “3” after the alphabet indicated the adhesive tape’s trademark Scotch[®] 3M super 33+ All the weather Vinyl Electrical Tape and YAZAKI P.V.C. Tape, respectively; the next two number indicated that the volunteer number.

Table 30 Concentration of extracted DNA from electrical adhesive tape (continued)

DNA samples	DNA concentration (ng/μl)	PCR efficiency ("0"= Ineffective, "1"= Effective)
ReSP101	1.0150	1
ReSP102	0.1805	1
ReSP103	0.2030	1
ReSP104	0.0705	0
ReSP105	0.6300	1
ReSP106	0.4545	1
ReSP107	0.1405	1
ReSP108	0.5100	1
ReSP109	0.3230	1
ReSP110	0.3045	1
ReSP201	0.1005	1
ReSP202	0.0840	0
ReSP203	0.0491	0
ReSP204	0.4290	1
ReSP205	0.2195	1
ReSP206	0.2275	1
ReSP207	1.4250	1
ReSP208	0.2520	1
ReSP209	0.4775	1
ReSP210	0.2925	1

“ReSP” indicated that the adhesive tape extracted by Phenol-Chloroform extraction method, the number “1” and “2” after the alphabet indicated the adhesive tape’s trademark Scotch[®] 3M 1710 Vinyl Electrical Tape and Scotch[®] 3M super 33+ All the weather Vinyl Electrical Tape, respectively; the next two number indicated that the volunteer number.

Table 30 Concentration of extracted DNA from electrical adhesive tape (continued)

DNA samples	DNA concentration (ng/μl)	PCR efficiency ("0"= Ineffective, "1"= Effective)
ReSP301	0.3325	1
ReSP302	0.1795	1
ReSP303	0.0231	0
ReSP304	0.3105	1
ReSP305	0.5900	1
ReSP306	0.2060	1
ReSP307	3.3650	1
ReSP308	0.1565	1
ReSP309	0.3745	1
ReSP310	0.0915	0

“ReSP” indicated that the adhesive tape extracted by Phenol-Chloroform extraction method, the number “3” after the alphabet indicated the adhesive tape’s trademark YAZAKI P.V.C. Tape, the next two number indicated that the volunteer number.

APPENDIX C

STR PROFILE OF THE REFERENCE DNA SAMPLE FROM THE VOLUNTEERS

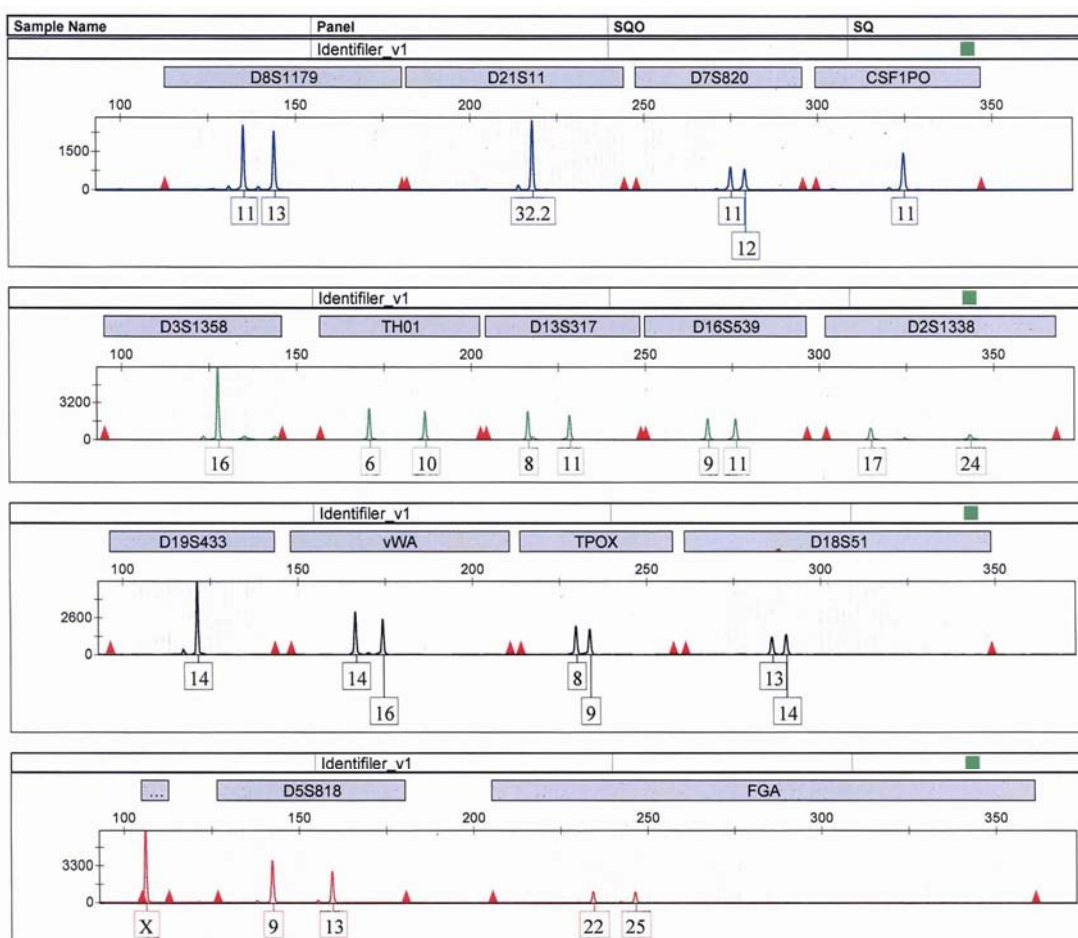


Figure 7 STR profile of the reference DNA sample from the volunteer no.1

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

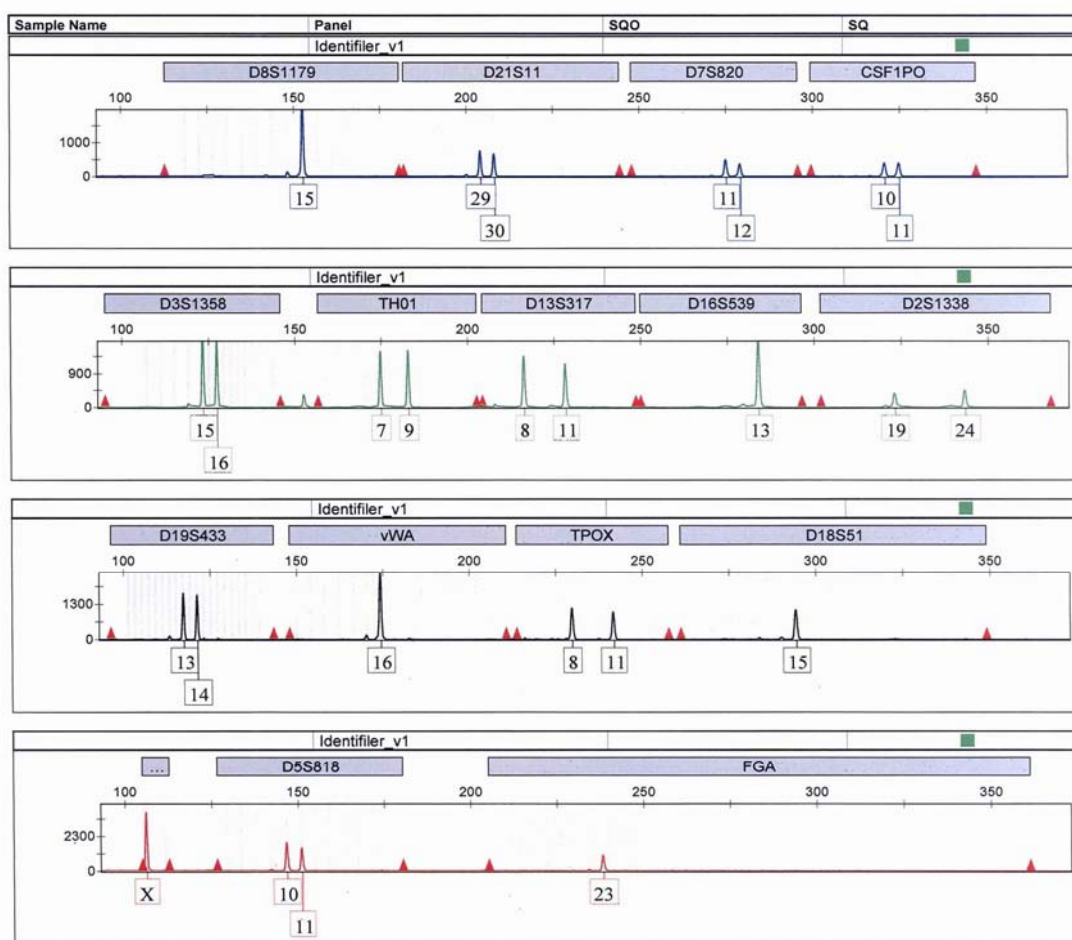


Figure 8 STR profile of the reference DNA sample from the volunteer no.2

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

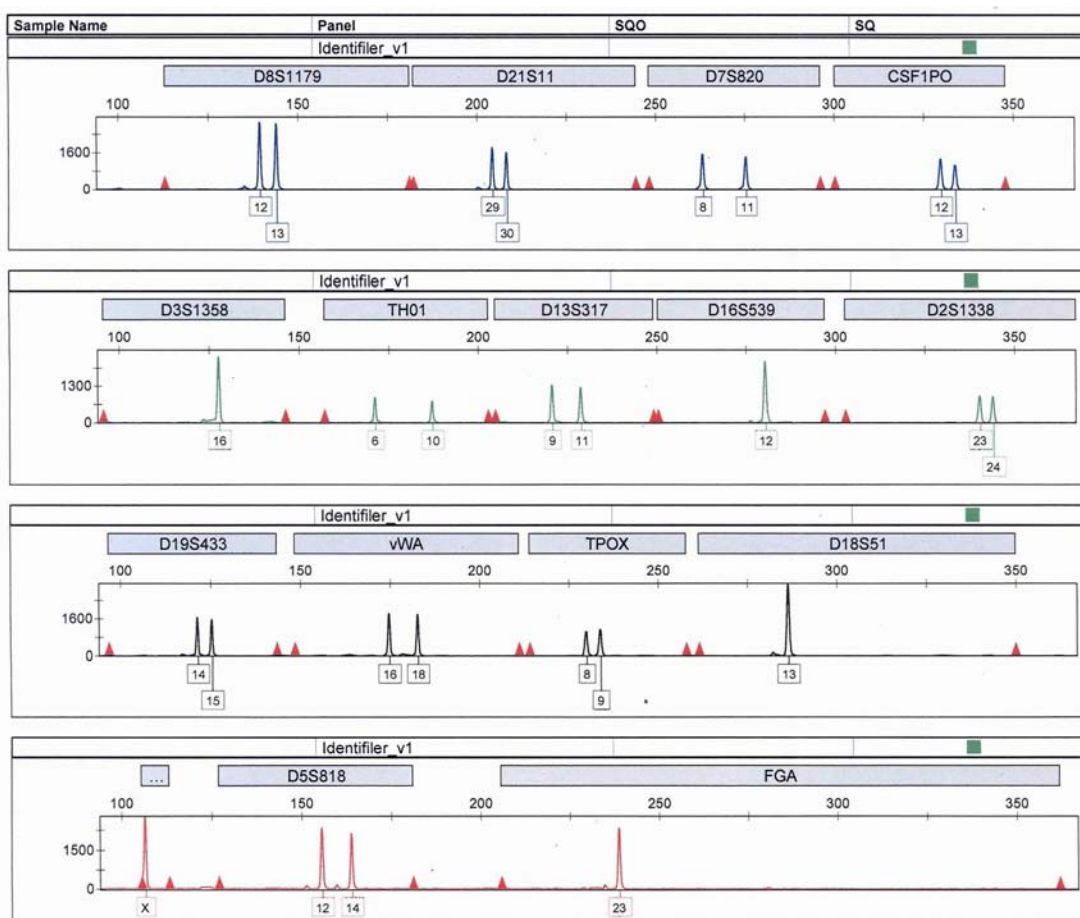


Figure 9 STR profile of the reference DNA sample from the volunteer no.3

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

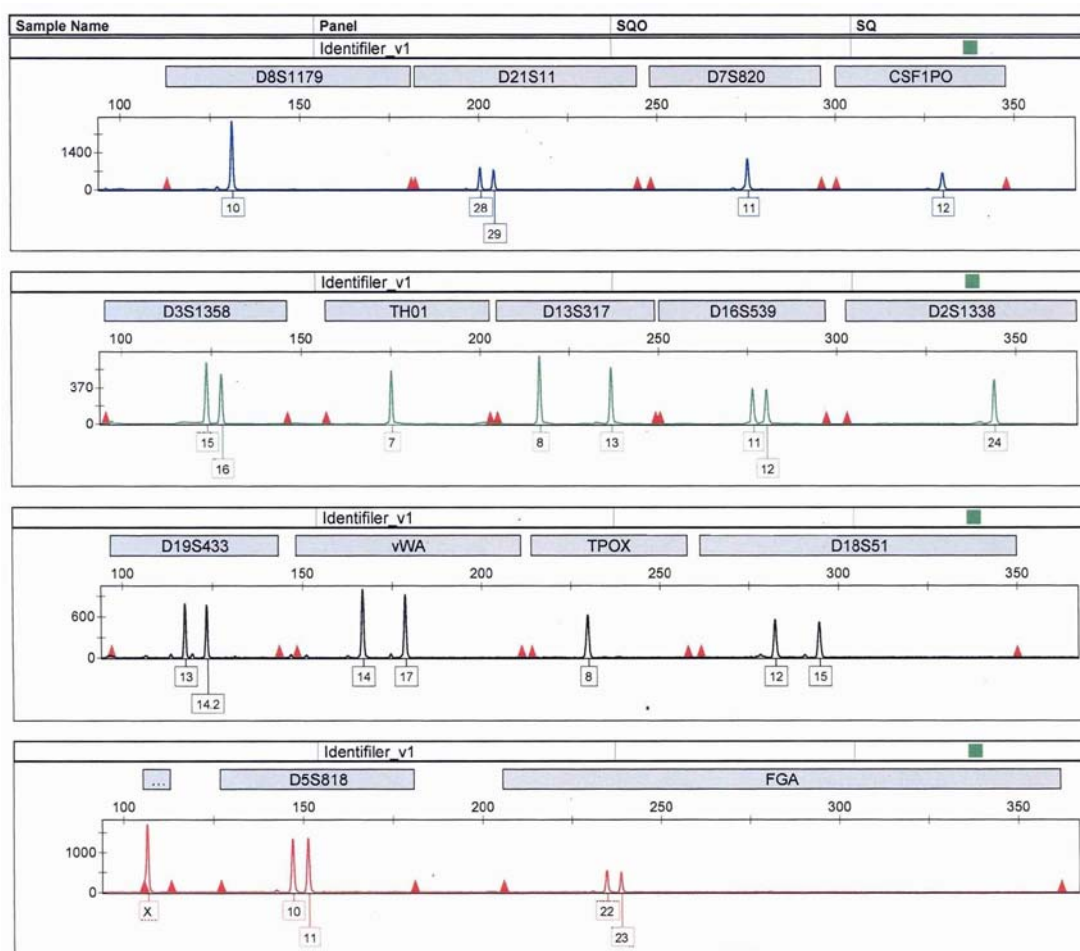


Figure 10 STR profile of the reference DNA sample from the volunteer no.4

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

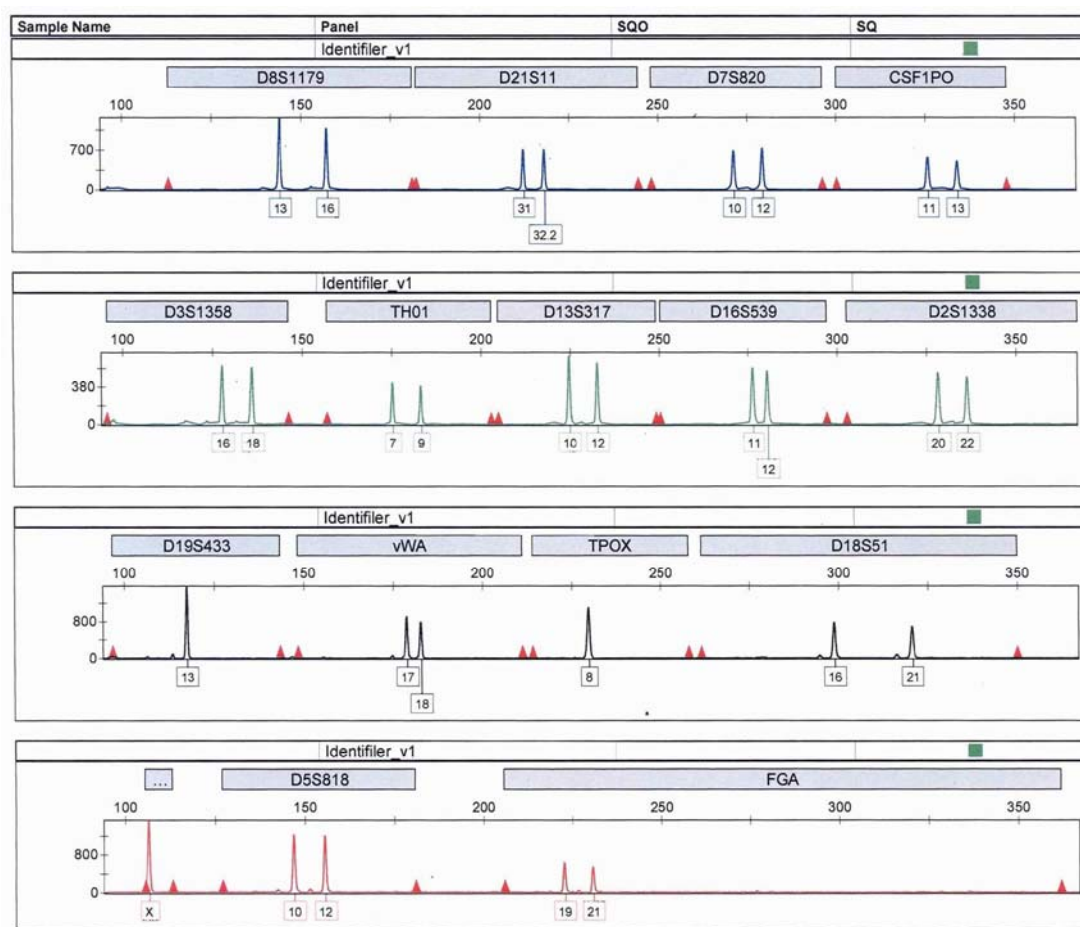


Figure 11 STR profile of the reference DNA sample from the volunteer no.5

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

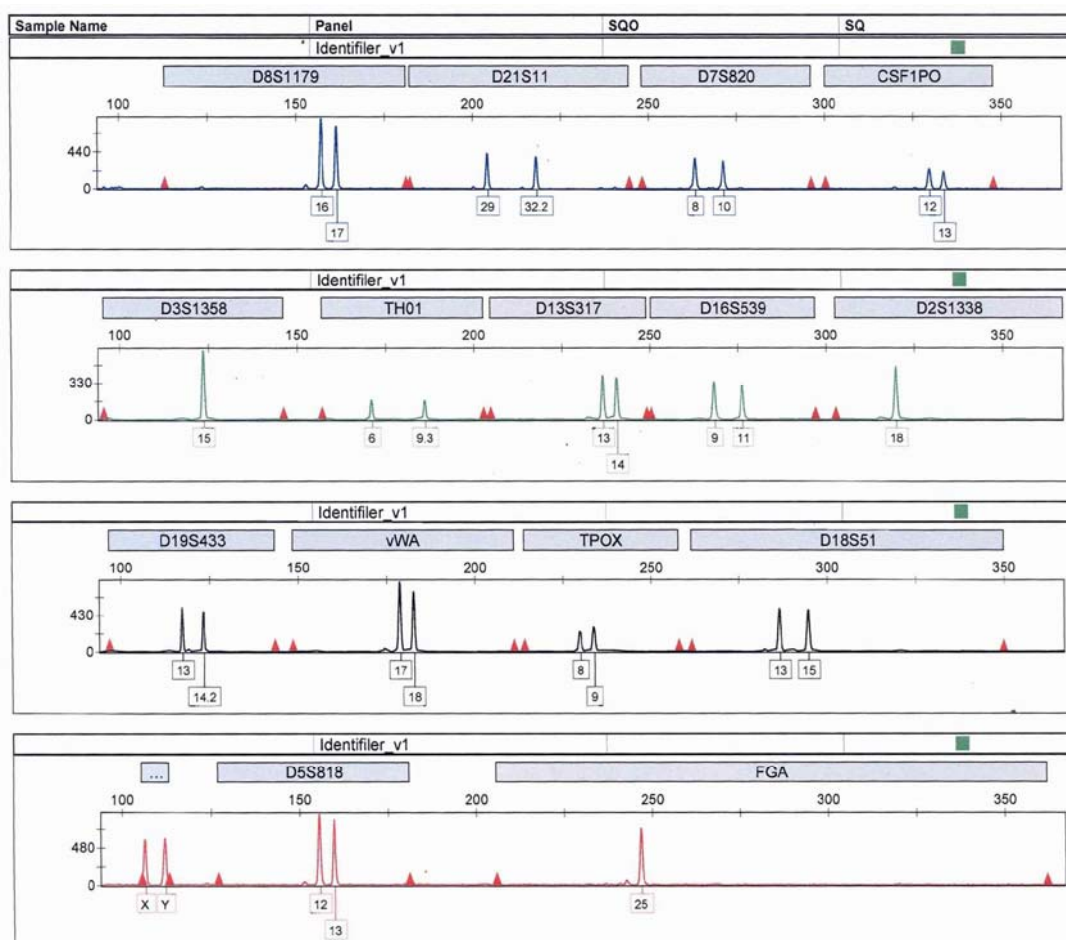


Figure 12 STR profile of the reference DNA sample from the volunteer no.6

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

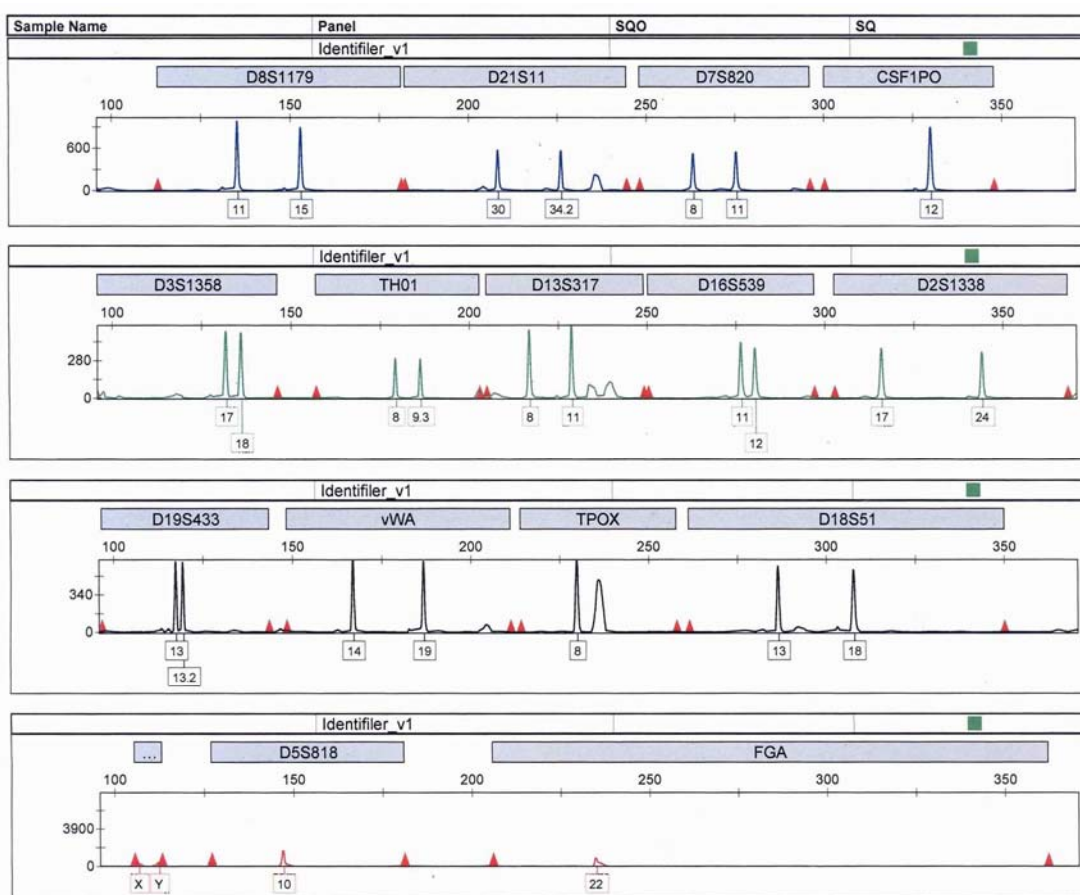


Figure 13 STR profile of the reference DNA sample from the volunteer no.7

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

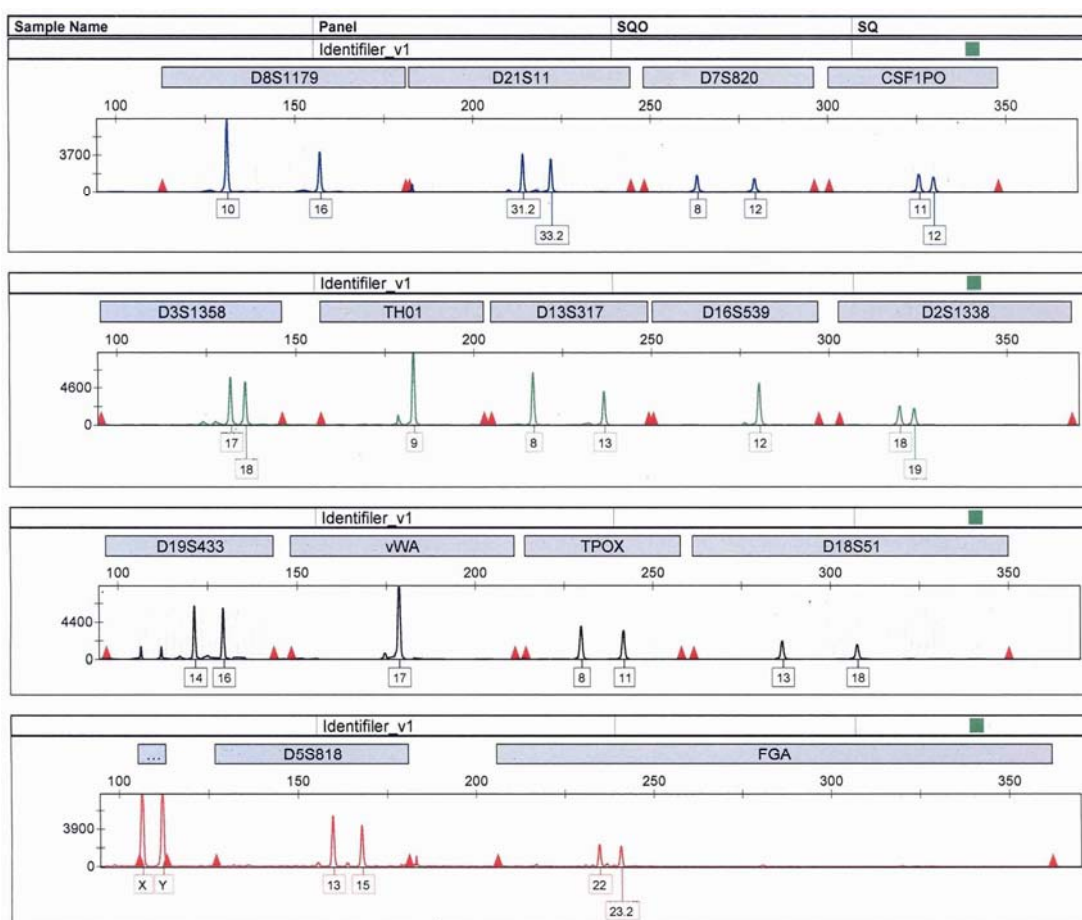


Figure 14 STR profile of the reference DNA sample from the volunteer no.8

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

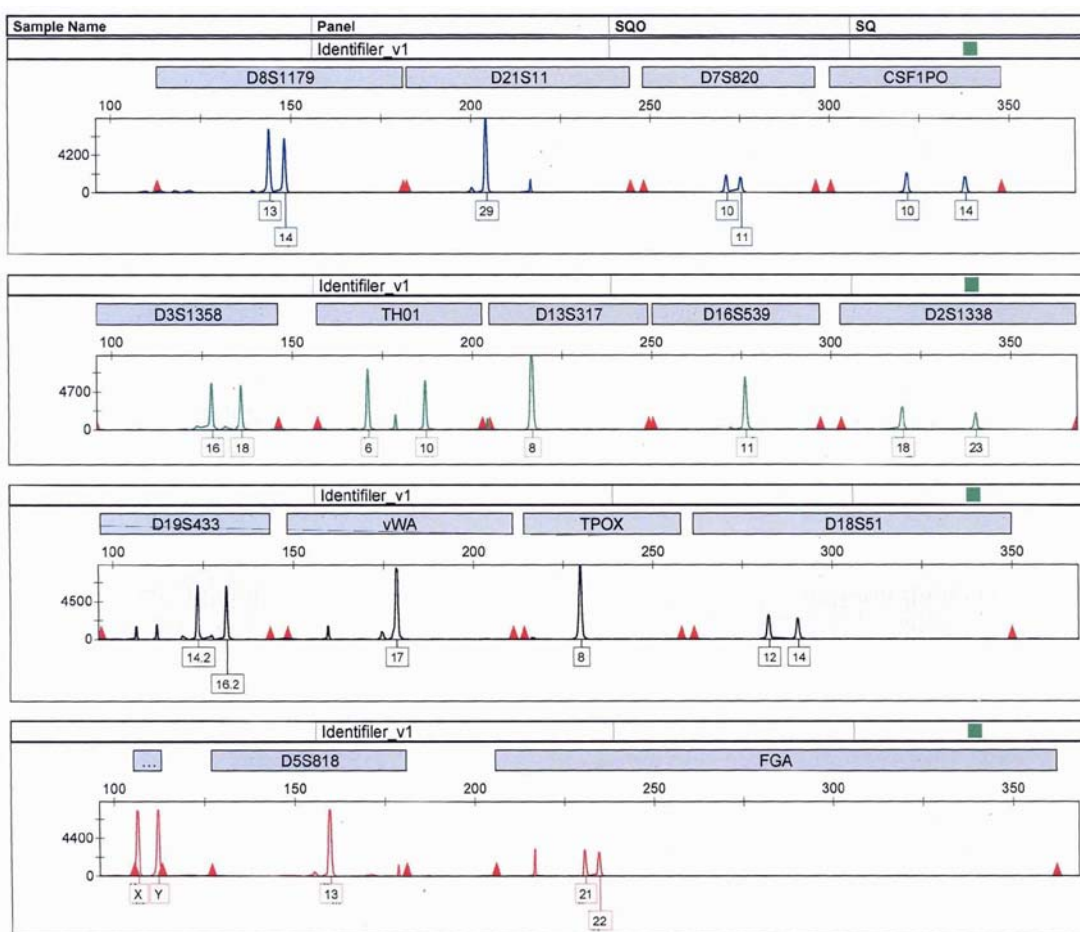


Figure 15 STR profile of the reference DNA sample from the volunteer no.9

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

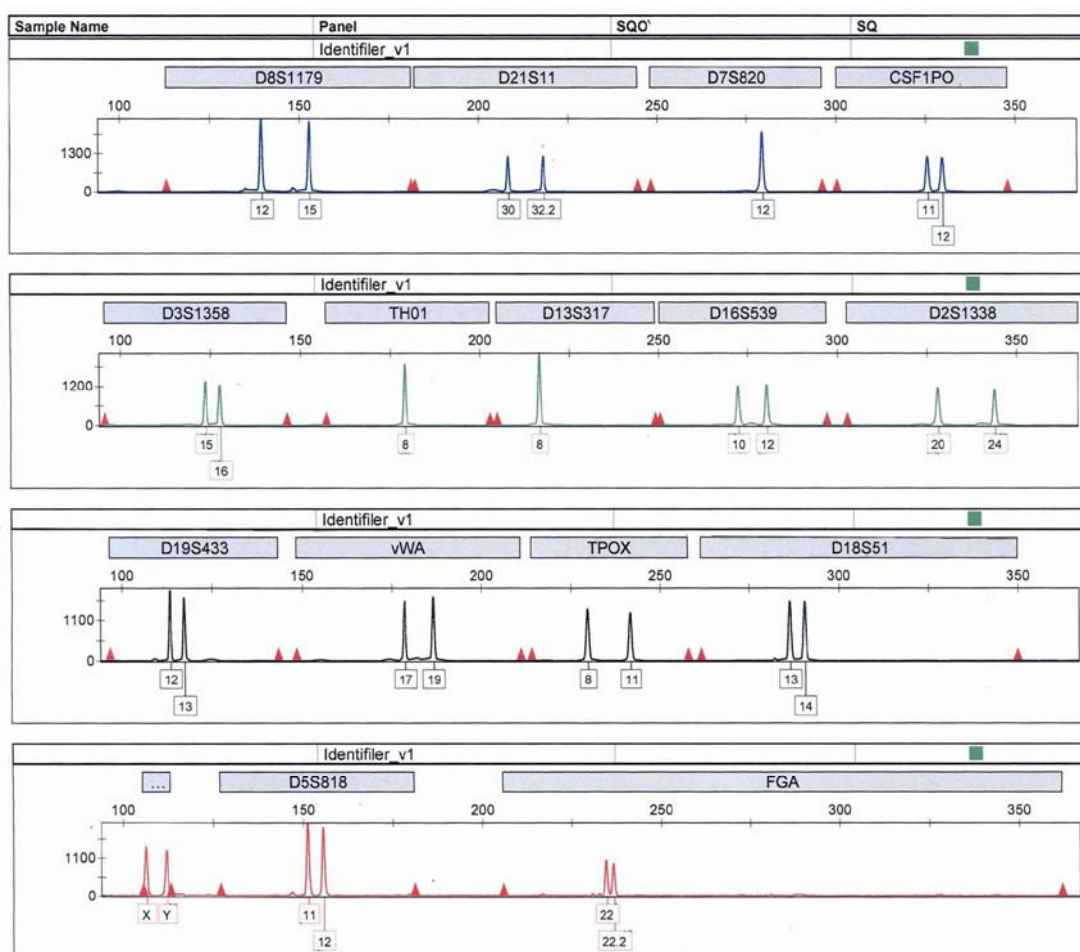


Figure 16 STR profile of the reference DNA sample from the volunteer no.10

Source : Ministry of Justice, Central Institute of Forensic Science, Forensic DNA Service Center, CIFS STAFF Reference DNA Profile (FM-DNA-051) (Bangkok : Forensic DNA Service Center, 2008), n. pag.

APPENDIX D

STR PROFILE OF THE DNA FROM EACH EXTRACTION METHOD

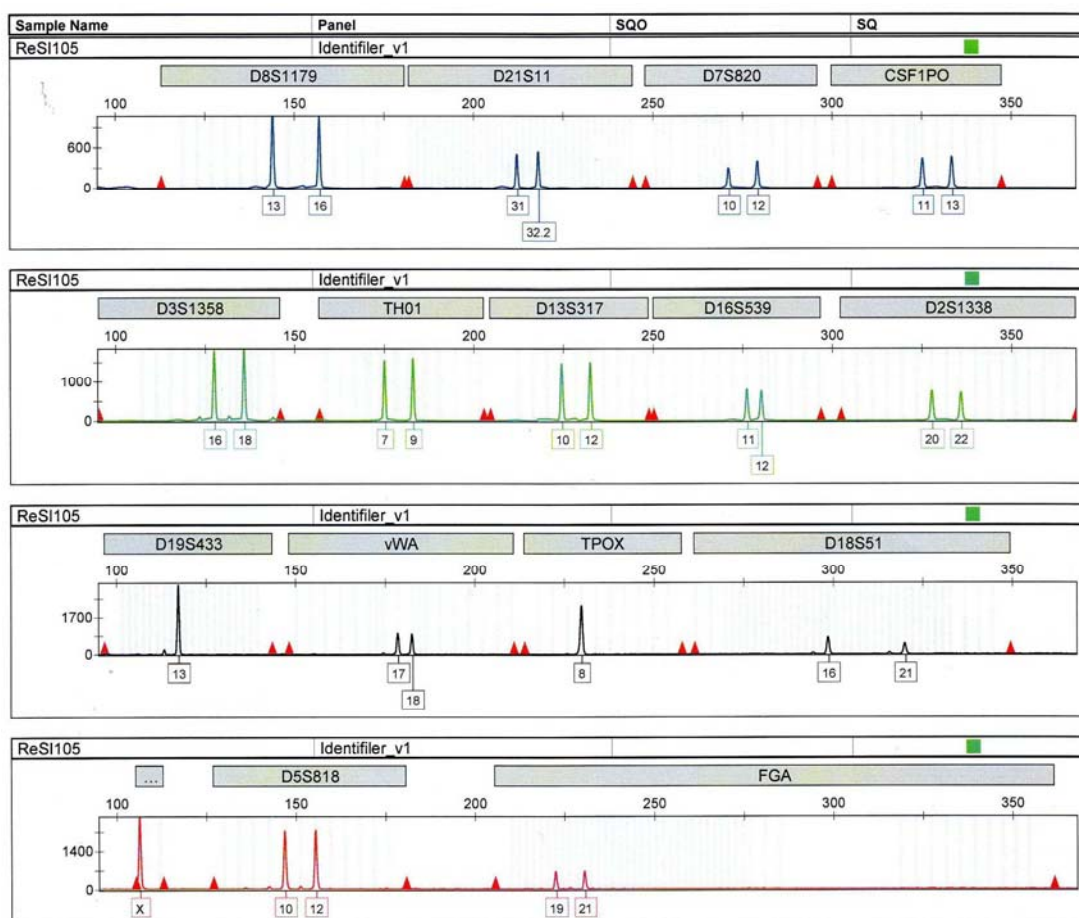


Figure 17 STR profile of the DNA from DNA IQ™ System extraction (ReSI) collected by Scotch® 3M 1710 Vinyl Electrical Tape of volunteer no.5

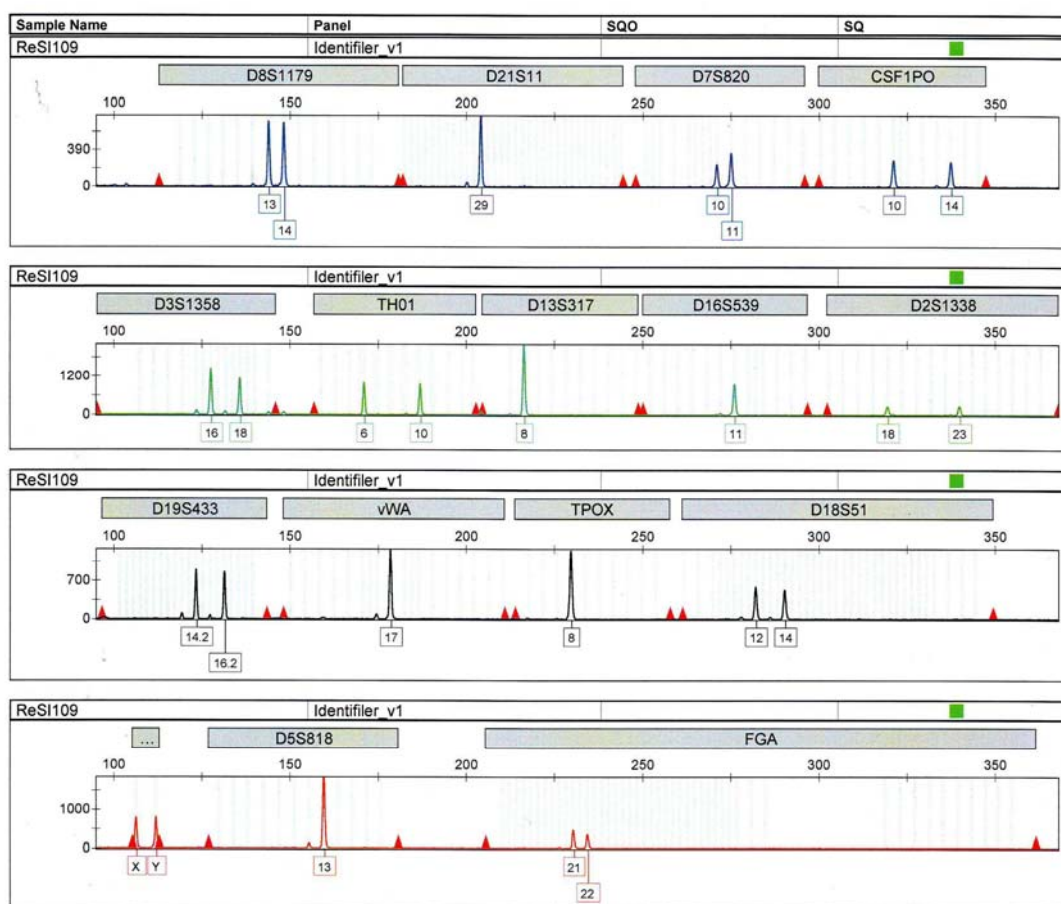


Figure 18 STR profile of the DNA from DNA IQ™ System extraction (ReSI) collected by Scotch® 3M 1710 Vinyl Electrical Tape of volunteer no.9

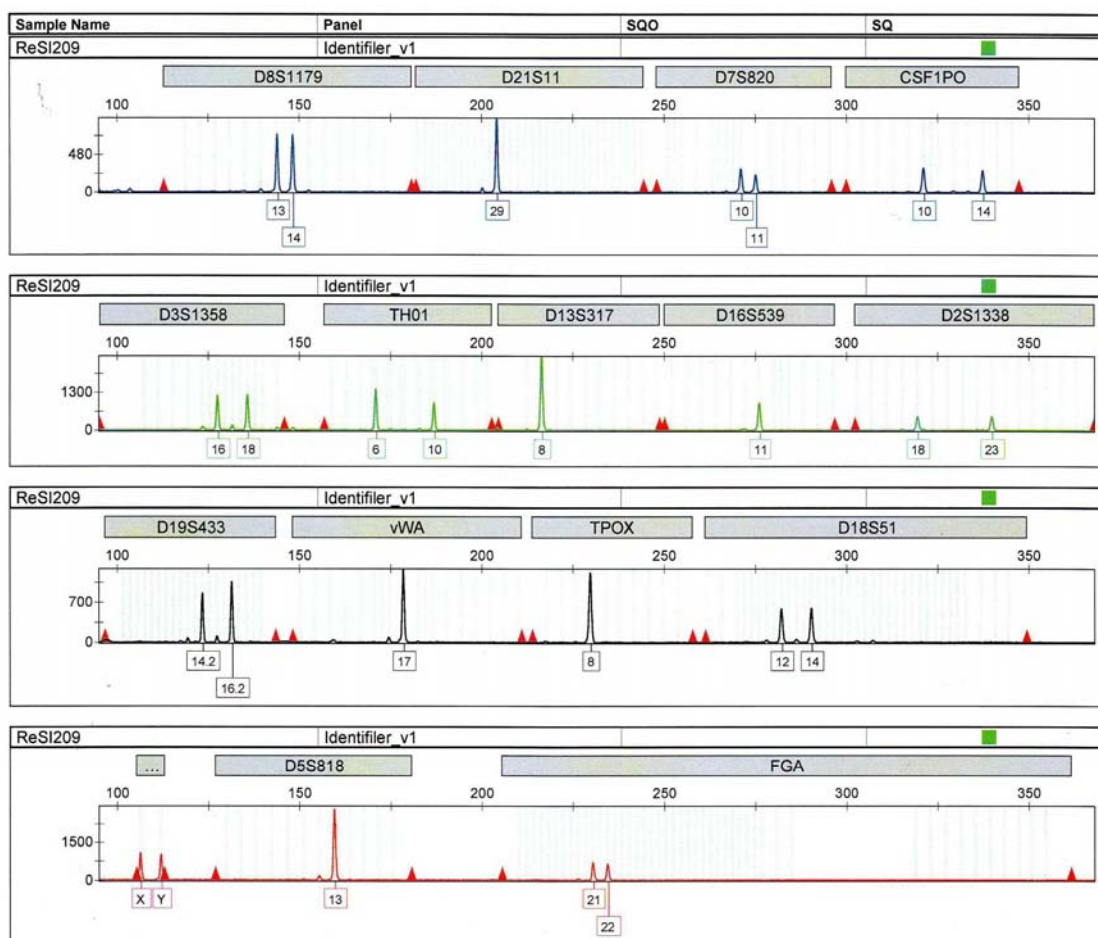


Figure 19 STR profile of the DNA from DNA IQ™ System extraction (ReSI) collected by Scotch® 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.9

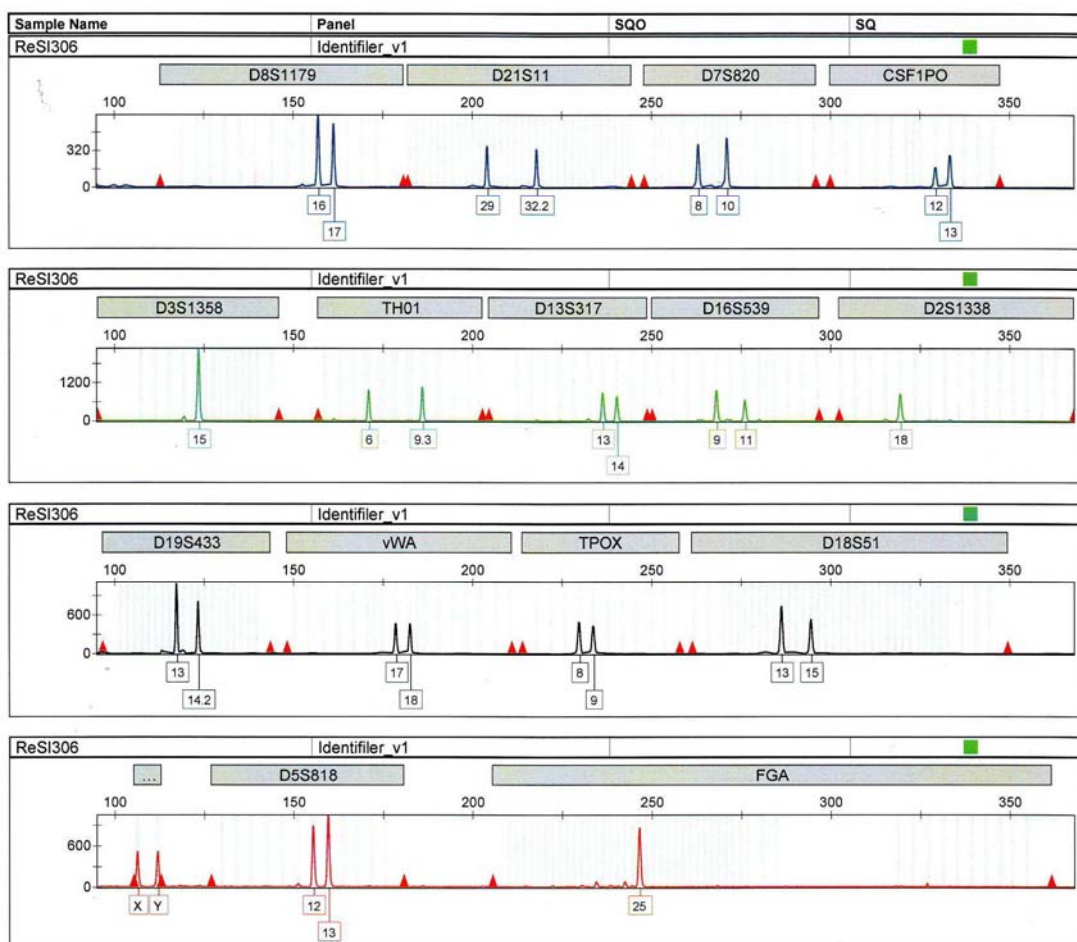


Figure 20 STR profile of the DNA from DNA IQ™ System extraction (ReSI) collected by YAZAKI P.V.C. Tape of volunteer no.6

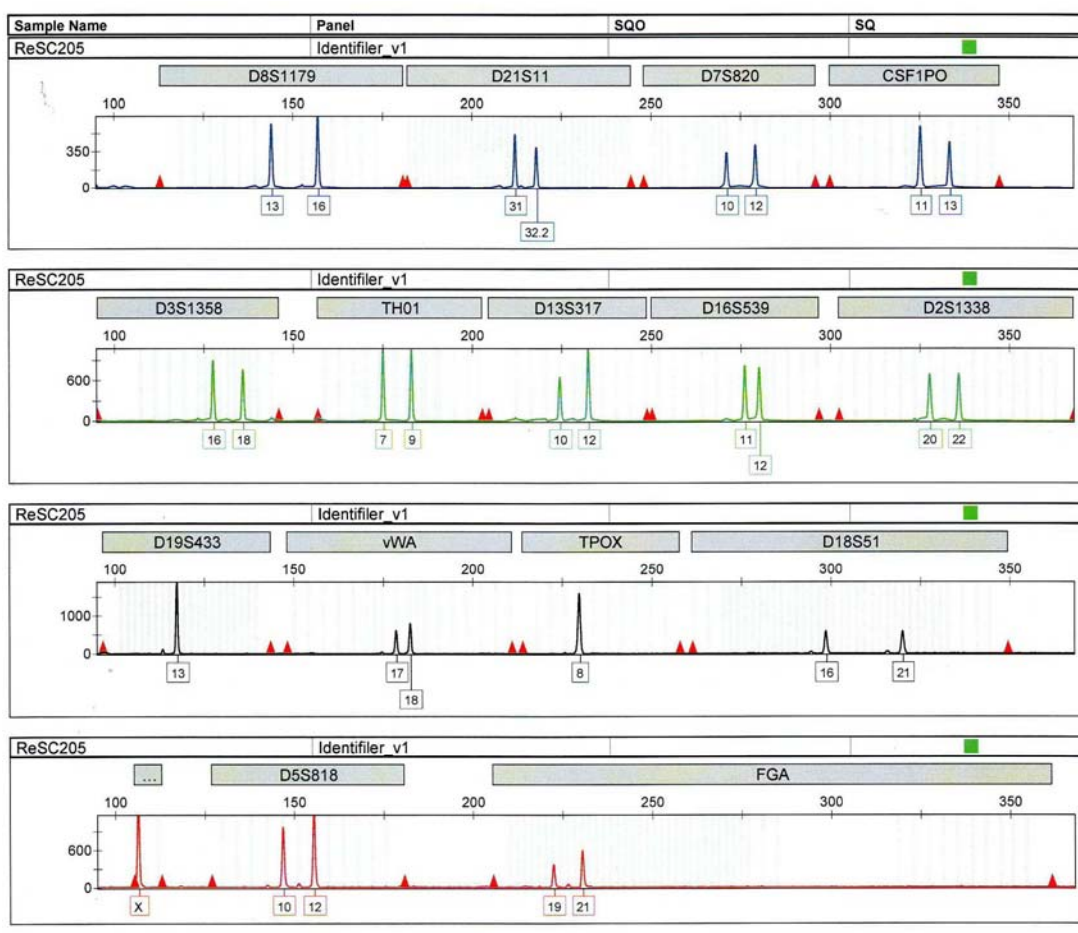


Figure 21 STR profile of the DNA from Chelex[®] 100 extraction (ReSC) collected by Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.5

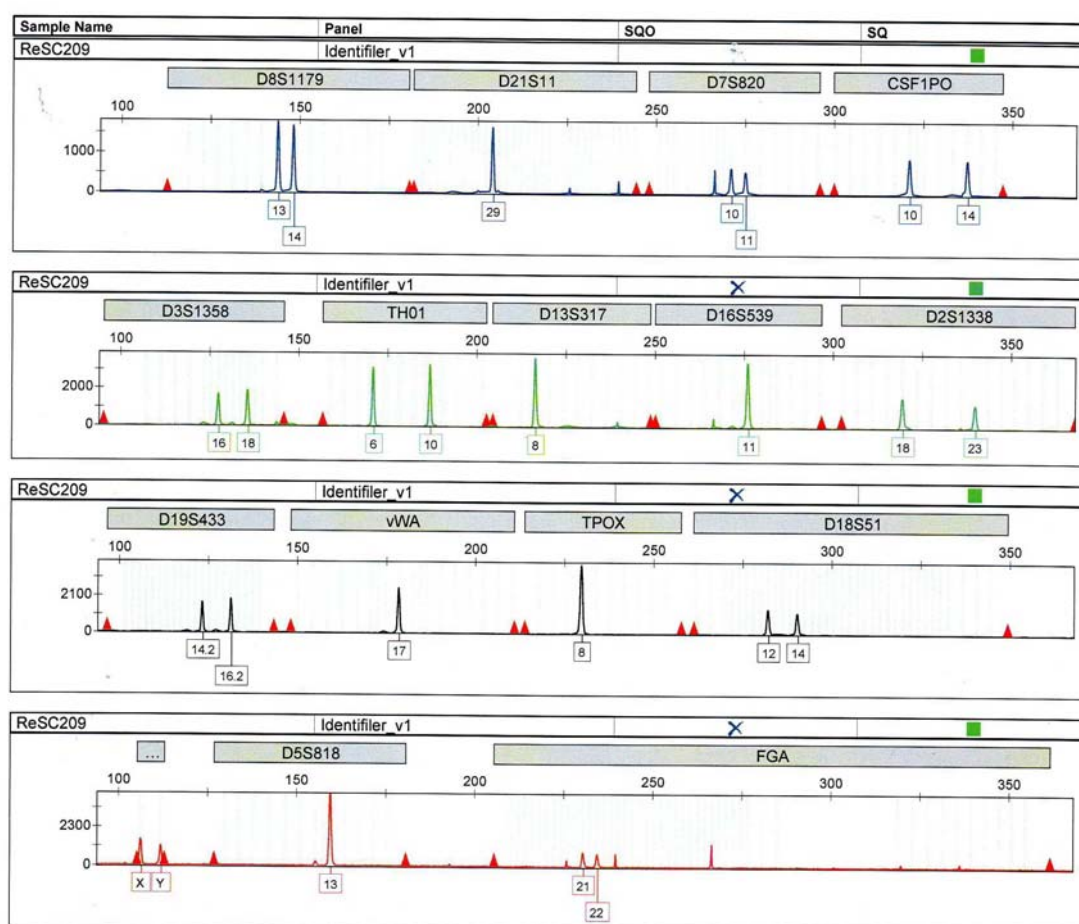


Figure 22 STR profile of the DNA from Chelex[®] 100 extraction (ReSC) collected by Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.9

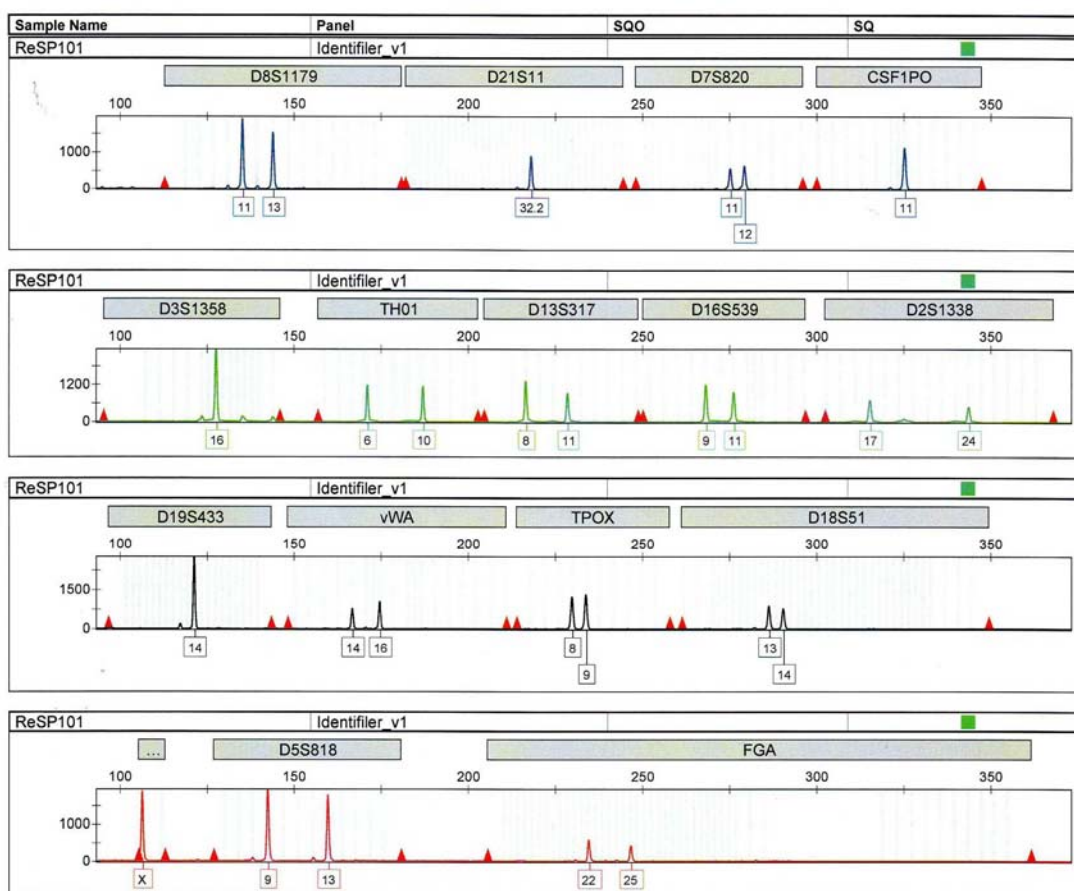


Figure 23 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by Scotch[®] 3M 1710 Vinyl Electrical Tape of volunteer no.1

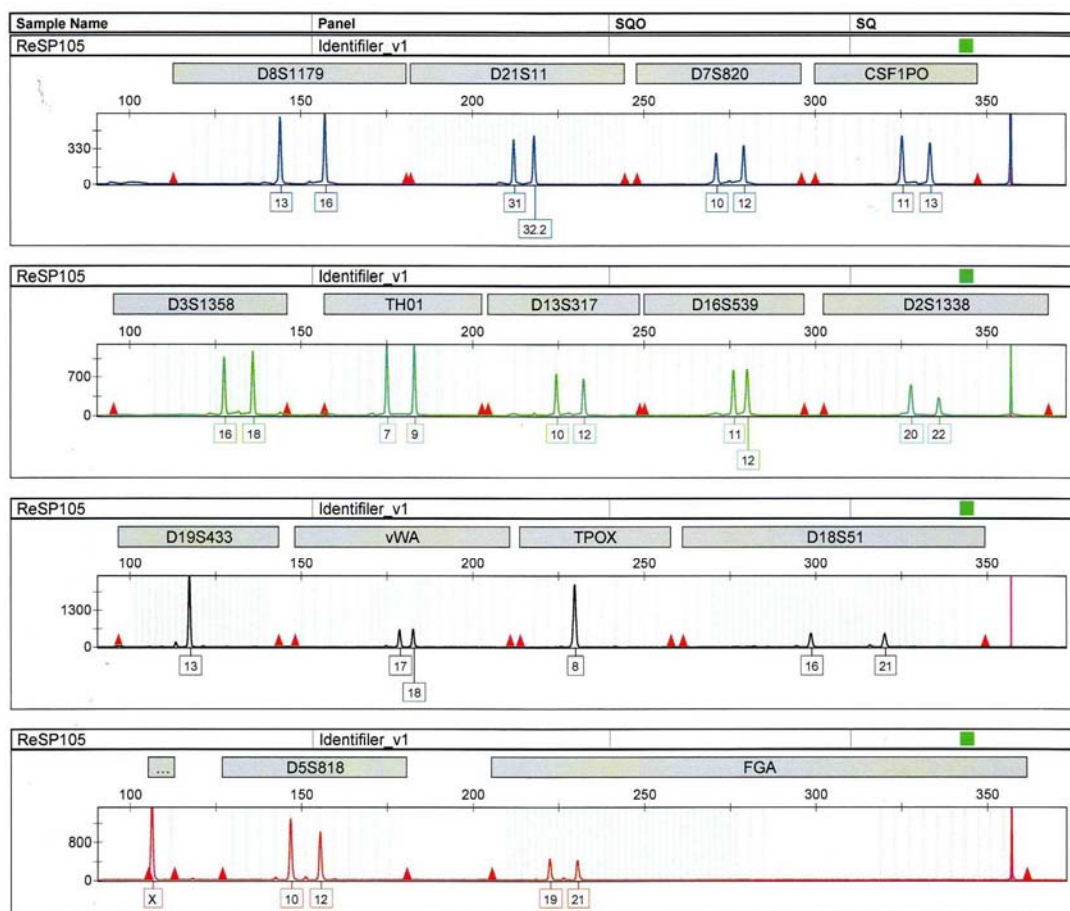


Figure 24 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by Scotch[®] 3M 1710 Vinyl Electrical Tape of volunteer no.5

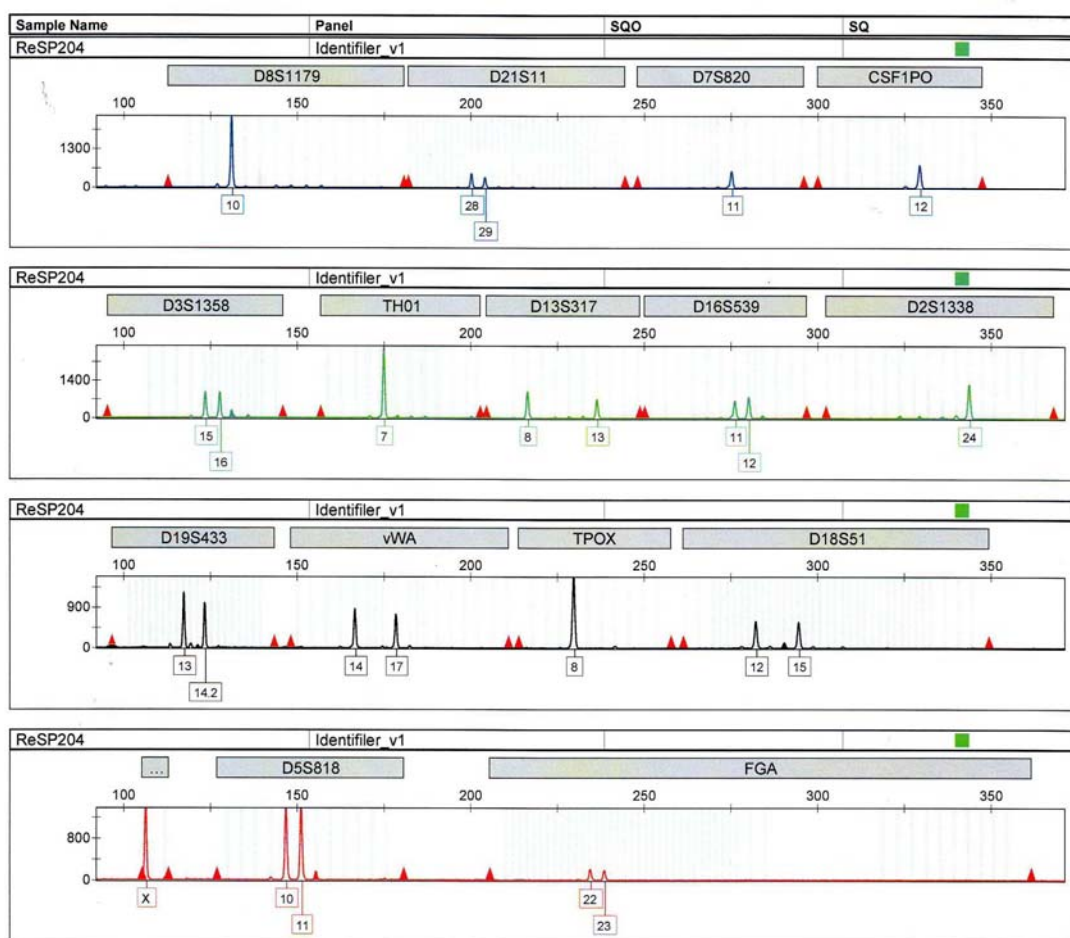


Figure 25 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.4

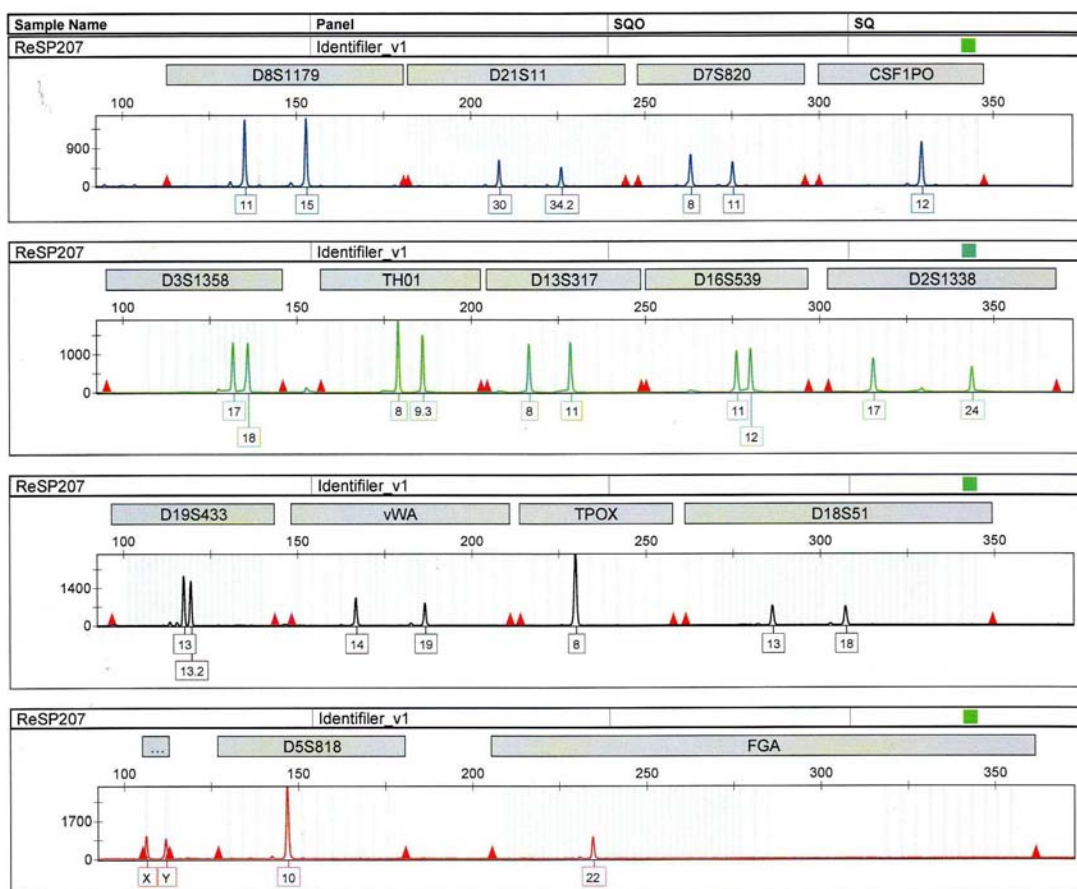


Figure 26 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by Scotch[®] 3M Super 33+ All Weather Vinyl Electrical Tape of volunteer no.7

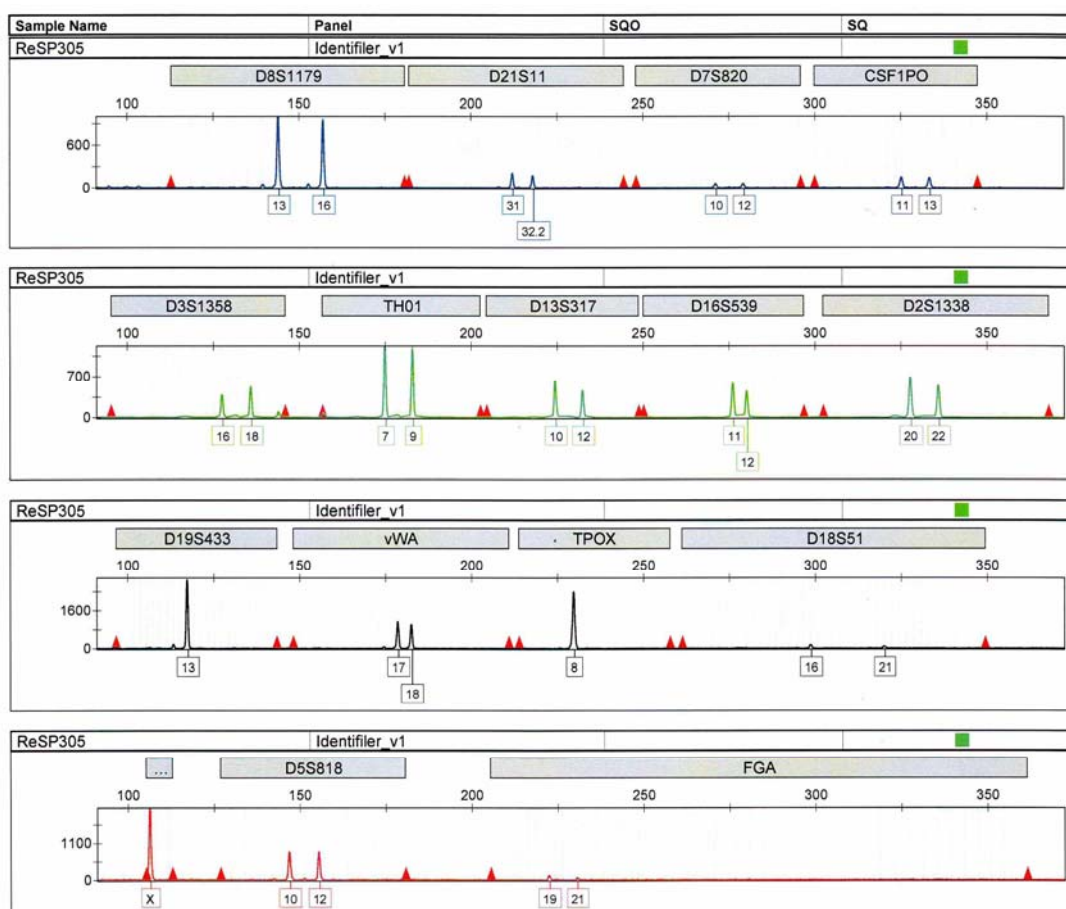


Figure 27 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by YAZAKI P.V.C. Tape of volunteer no.5

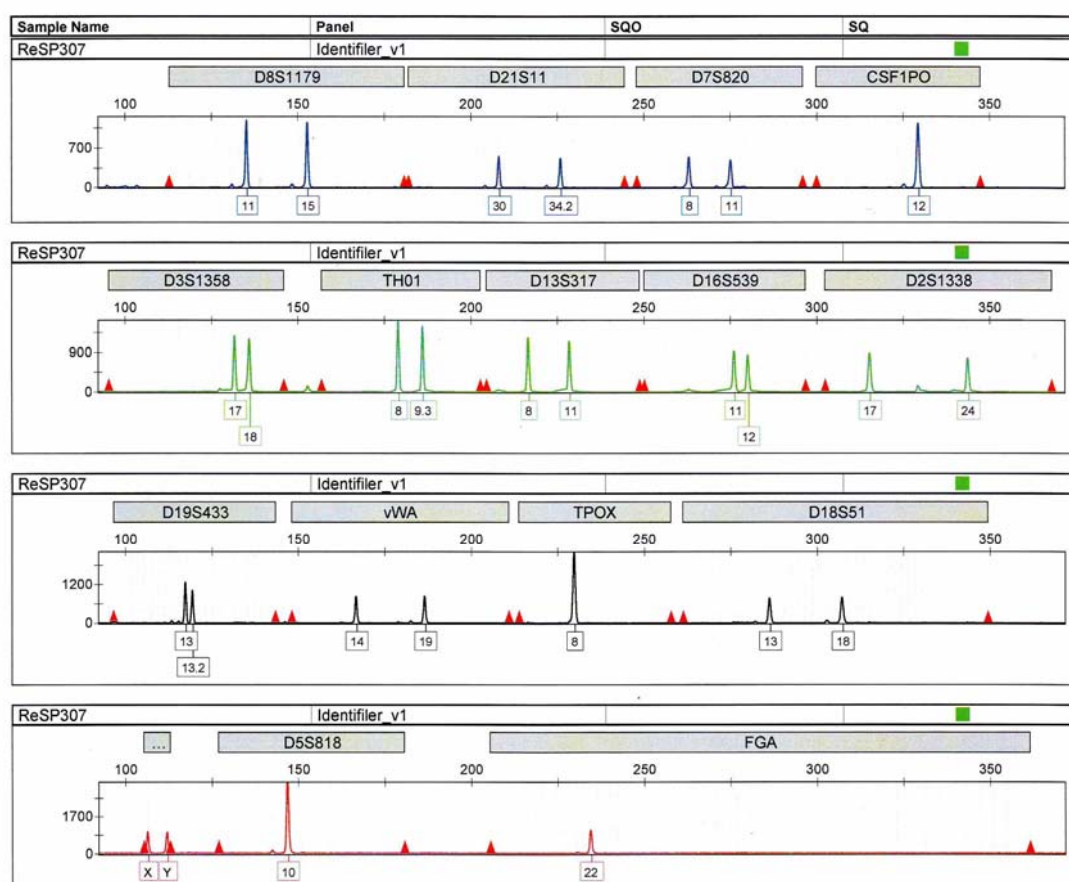


Figure 28 STR profile of the DNA from Phenol: Chloroform extraction (ReSP) collected by YAZAKI P.V.C. Tape of volunteer no.7

APPENDIX E
SUGGESTED PROTOCOL

Suggested protocol for DNA extraction from electrical adhesive tape by DNA IQ™ System

1. Place the electrical adhesive tape in sterile 50 ml centrifuge tube
2. Add 4 ml of prepared Lysis Buffer. Close the lid, and incubate the tube overnight room temperature.
3. Vortex tube for 10 seconds at high speed, and incubate at 95 °C for 30 minutes.
4. Remove the tube from the heat source, and transfer the solution to a new 15 ml centrifuge tube
5. Vortex the stock resin bottle for 10 seconds at high speed or until resin is thoroughly mixed. Add 7 µl of DNA IQ. Resin to the sample. Keep the stock resin resuspended while dispensing to obtain uniform results.
6. Vortex the sample/Lysis Buffer/resin mixture for 3 seconds at high speed. Incubate at room temperature for 5 minutes. Vortex mixture for 3 seconds once every minute during this 5 minute incubation.
7. Vortex tube for 2 seconds at high speed. Place tube in the magnetic stand. Separation will occur instantly.

Note: If resin does not form a distinct pellet on the side of the tube, vortex the tube and quickly place back in the stand.
8. Discard the solution until indicate volume 1 ml, vortex for 2 seconds and place the tube in the magnetic stand. Then remove all of the solution to a new 1.5 ml microcentrifuge tube and place tube in the magnetic stand.
9. Carefully remove and discard all of the solution without disturbing the resin pellet on the side of the tube.

Note: If some resin is drawn up in tip, gently expel resin back into tube to allow re-separation.
10. Add 100 µl of prepared Lysis Butter. Remove the tube from the magnetic stand, and vortex for 2 seconds at high speed.
11. Return tube to the magnetic stand, and discard all Lysis Buffer.
12. Add 100 µl of prepared 1X Wash Buffer. Remove tube from the magnetic stand, and vortex for 2 seconds at high speed.
13. Return tube to the magnetic stand, and discard all Wash Buffer.

14. Repeat Steps 12 and 13 two more times for a total of three washes. Be sure that all of the solution has been removed after the last wash.

15. With the tube in the magnetic stand and the lid open, air-dry the resin for 5 minutes.

Note: Do not dry for more than 20 minutes, as this may inhibit removal of DNA.

16. Add 15 μ l of Elution Buffer.

17. Close the lid and vortex the tube for 2 seconds at high speed. Incubate the tube at 65 °C for 5 minutes.

18. Remove the tube from the heat source, and vortex for 2 seconds at high speed.

Immediately place the tube on the magnetic stand.

Note: Tubes must remain hot until placed in the magnetic stand or yield will decrease.

19. Carefully transfer the DNA-containing solution to a container of choice.

Note: DNA can be stored at 4 °C for short-term storage or at -20 or -70 °C for long-term storage.

Suggested protocol for DNA concentration by Microcon™ YM-100

1. Assemble and label a fresh microcon unit for each sample. Add 50 µl of TE buffer (0.1 mM EDTA) the filter side of the membrane (upper side).

Note: The maximum volume Microcon 100's (blue retanate cup) can accept is 500 µl.
2. Add sample to corresponding microcon retanate cup.
3. Centrifuge microcon with lid closed for 5 minutes at 8,000 rpm until entire volume is filtered.
4. Add 100 µl TE buffer and 200 µl of extract DNA to filter side of cup, and centrifuge for 5 minutes at 8,000 rpm until the entire volume is filtered. Discard filtrate, return cup to microcon tube.
5. Label new microcon tube for each sample. Carefully loosen cup in original microcon tube but do not remove. Add 20 µl of TE buffer to the filter side of cup of original microcon unit. Vortex tube with new labeled microcon tube held over top, and centrifuge for 5 minutes at 10,000 rpm.
6. Discard retanate cup, retain filtrate, and store appropriately.
7. Pooling samples. If necessary to pool samples, prepare new microcon unit and add 50 µl of TE buffer to the filter side of the membrane (upper side). Pool the 20 µl volumes from several tubes of the same sample into microcon retanate cup. Centrifuge microcon for at least 5 minutes at 10,000 rpm with lid closed until the entire volume is filtered. Discard filtrate, return cup to microcon tube. Elute DNA from membrane following Steps 5-6.

BIOGRAPHY

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2008	Master of Science, Program of Forensic Science, Graduate School, Silpakorn University
Working Experience	
2008 - present	Staff in the project of convicted felons DNA database, Forensic DNA Service Center, Central Institute of Forensic Science, Ministry of Justice