Identification of Murder Victims' Cranial Bone Mutilation Using Forensic Medicine, Anthropology, and Genotype DNA Approaches

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ABSTRACT

The objective of forensic identification is to aid investigators in ascertaining an individual's identity. Reliability in identifying human remains from natural catastrophes, man-made disasters, and situations involving missing individuals has significantly advanced. Anthropological examination and forensic genetic profiling are particularly beneficial when traditional identification procedures, such as fingerprinting or ocular recognition, are not feasible. Even in cases where only skeletal remains are present, anthropological inquiries and forensic genetic examination of the remaining body parts can ascertain the identity and familial connection of the surviving individual. We have found a decapitated skull that is missing its lower jaw and comprises several cervical bones. Investigators think that the decapitated head is a component of a victim who was previously interred. We conducted an autopsy, an anthropological analysis and report the process of identifying skull bone mutilations in murder victims using forensic medicine, anthropology, and DNA genotyping approaches. We performed autopsy and anthropological investigations to collect data from the skeletal remains and a genetic analysis by collecting tooth and blood samples from victims' parents. These samples were then utilized for DNA extraction, calculation of DNA rate and purity, amplification, and identification of genotype. After investigation, we discovered a single cranial bone and four cervical bones. Under macroscopic examination, the bones exhibit a striking resemblance to the structure of a human head and neck. The presence of tissue still attached to the bones indicates that the time of death exceeds 10 days. The complete destruction of the cranial bones indicates that they belong to individuals aged between 21 and 39 years. The presence of shovel-shaped teeth, a rounded palatal form, straight palatal sutures, and molar teeth with four cusps provide strong identification of the deceased individual as belonging to the Mongoloid race. The assessment of height is challenging due to the absence lengthy bones

Keyword: Antrophology, Autopsy, Genetic Profiling, Identification, Skeletal Remains.

INTRODUCTION

The objective of forensic identification is to aid investigators in ascertaining an individual's identify.1 Significant advancements have been achieved in reliably identifying human remains from natural catastrophes, man-made disasters, and situations involving missing individuals.² Anthropological examination and forensic genetic profiling are particularly beneficial when traditional identification procedures such as fingerprinting or ocular recognition are not feasible. Even in cases when only skeletal remains are present, anthropological inquiries and forensic genetic examination of the remaining body parts can ascertain the identify and familial connection of the surviving individual.3

We have found a decapitated skull that is missing its lower jaw and comprises several cervical bones. Investigators think that the decapitated head is a component of a victim who was previously interred. A decapitated human corpse was found one week ago. The Mobile Automatic Multi Biometric Identification System (MAMBIS) technology effectively recognized the deceased individual at the location of the crime. Subsequently, his family interred the deceased's remains, adamantly declining to conduct an autopsy. We conducted an analysis of the head parts.

METHODS

We conducted an autopsy, an anthropological analysis, and a DNA profiling investigation. We performed autopsy and anthropological investigations to collect data from the skeletal remains. This involved ascertaining the human provenance of the skull, establishing whether it came from a single individual or several individuals, calculating the time since death, defining the gender, finding any indications of violence, diagnosing the reason of death, and determining the manner of death. We performed a genetic analysis by collecting tooth and blood samples from the alleged parents of the victim (Table 1). These samples were then utilized for DNA extraction, calculation of DNA rate and purity, amplification, and identification of genotype.

Extraction of DNA

The three samples were obtained by utilizing the 5% chelex procedure, which involves combining 5 grams of chelex with 100 ml of a pH 8.0 TE buffer solution. The TE Buffer Solution is prepared by dissolving 0.61 g of Tris Base and 0.0186 g of EDTA in 500 ml of ddH20. Subsequently, introduce the

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mixed solution into either a 1 M HCL or 1 M NaOH solution until the pH of the resulting solution reaches 8.0. Prior to usage, thoroughly mix the 5% chelex to ensure that there are no residual deposits in the solution. For sample X1, include 400 μ l of Chelex 5%, 2 μ l of prot K with a concentration of 10 mg/ml, and 18 μ l of DTT with a concentration of 0.05M. Meanwhile, samples X2 and X3 introduced 400 μ l of Chalex 5% and 10 μ l of Prot K 10 mg/ml into the sample tube. The X1 sample tubes were subjected to incubation in a water bath with shaking at a temperature of 56.0 °C for a duration of 4 hours, while the X2 and X3 sample tubes were incubated for 1 hour. Subsequently, we subjected each sample to vortexing for a duration of 10 seconds and centrifugation at a speed of 13,000 revolutions per minute for a period of 3 minutes. The supernatants obtained from the tubes were transferred into individual 15-ml tubes and kept at a temperature of 40 °C for further analysis.

Quantification of DNA

 $0.5\,\mu l$ of human standard primer, $12.5\,\mu l$ of PCR reaction from the ABI Human Quantifiler kit from Applied Biosystems in the USA, and $2\,\mu l$ of sample were added to a 96-well plate also made by Applied Biosystems in the USA. Subsequently, we sealed the plate, subjected it to vortexing, and spun it at a speed of 3,000 revolutions per minute for a duration of 1 minute. We performed DNA quantification by placing the sample plate into the ABI 7,500 Real-Time PCR machine manufactured by Applied Biosystems in the United States.

DNA amplification

Thisstudy involves the amplification of DNA samples using a combination of 21 microsatellite encoding markers, one Y-chromosome encoding marker, one Y-indel marker, and one sex marker (Amelogenin). These markers are put together in the GlobalFiler^{*} kit, which is manufactured by Applied Biosystems in the United States. Table 1. The PCR tube was filled with 7.5 μ l of matrix nix, 2.5 μ l of forward and reverse primer pairs, and 1 ng of the extracted DNA sample. Subsequently, we introduced the PCR tube, which held the sample, into the GeneAmp PCR System 9700 manufactured by Applied Biosystems in the United States. The machine's amplification process consists of many phases. It begins with an initial heating step at a temperature of 95°C for a duration of 1 minute. This is followed by 96 cycles, each consisting of denaturation at 94°C for 10 seconds, attachment at 59°C for 90 seconds, extension at 60°C for 10 minutes, and a final extension step at 4°C for a period of 24 hours.

Microsatellites

We loaded a 96-well plate (Applied Biosystem, USA) with a combined volume of 9.6 liters of HiDi formamide, 0.4 liters of 600 LIZ, and 1 liter of PCR samples. Subsequently, we hermetically closed the plate and subjected it to centrifugation at a speed of 3,000 revolutions per minute for a duration of 1 minute. The sample was subjected to a temperature of 950 °C for a duration of 3 minutes, followed by a further cooling process in a freezer at -250 °C for 3 minutes. The samples were subsequently evaluated using the ABI 3500XL Genetic Analyzer, manufactured by Applied Biosystems in the United States.

Genotype Identification

The data obtained from the ABI 3500XL Genetic Analyzer (Applied Biosystems, USA) was processed using GeneMapper* ID-X Software v1.4 (Applied Biosystems, USA)

No	Type of Evidence	Lab Code	Information
1.	Teeth	24058_1	X1
2.	Mother's blood sample	24058_2	X2
3.	Father's blood sample	24058_3	X3



Figure 1. Bone examination. The examination focuses on the cervical bones 1-4 (a), skull (b), and upper jaw (c).

RESULTS

Autopsy and Anthropologic Findings

After careful investigation, we discovered a single cranial bone and four cervical bones. Under macroscopic examination, the bones exhibit a striking resemblance to the structure of a human head and neck (Figure 1). They originate from a singular individual. The presence of tissue still attached to the bones indicates that the time of death exceeds 10 days. Furthermore, the anatomical features suggest a male gender, as evidenced by the sloping shape of the forehead bone, the protrusion of the glabella, the roughly square shape of the orbital cavity, and the noticeable prominence of the external occipital protuberance. The complete destruction of the cranial bones indicates that they belong to individuals aged between 21 and 39 years. The presence of shovelshaped teeth, a rounded palatal form, straight palatal sutures, and molar teeth with four cusps provide strong identification of the deceased individual as belonging to the Mongoloid race. The assessment of height is challenging due to the absence of lengthy bones. However, there were indications of violence, such as blood seepage at the back of the head and the base of the 4th cervical vertebrae. The cause of death was determined to be a sharp force applied to the neck, and the mode of death was classified as unnatural.

DNA Findings

DNA examination showed the following results. The DNA profile labeled as 24058_1 is a partial match to the DNA profile labeled as 24058_2, indicating that X2 is the biological offspring of the X1. The DNA profile 14058_1, which is half of lab code 24008_3, confirms that X3 is the biological of X1. The paternal index is 1: 408.064,538, indicating a 99.999% probability (Table 2).

DISCUSSION

Identifying human remains is crucial for both legal and humanitarian purposes.⁴ The forensic identification of human remains is a legal process where the jurisdictional authority confirms the identity of the deceased by comparing information on missing individuals with the characteristics of the unidentified remains. This confirmation is officially documented by the authority's signature on a death certificate.

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Table 2. Comparison of the victim's DNA profile with the alleged victim's parents.											
Lab Code	D8S1179	D21S11	D7S820	CSF1PO	D3S1358	TH01	D13S317	D16s539			
24058_1	12 13	29 30.2	8 8	11 11	16 17	6 9	9 11	10 13			
24058_2	11 12	29 30.2	8 11	10 11	16 17	69	10 11	11 13			
24058_3	13 15	29 31	8 8	11 12	15 16	7 9	9 14	10 11			
Lab Code	D2S1338	D19S433	vWA	TPOX	D18S51	AMEL	D5S818	FGA			
24058_1	16 20	13 14	17 18	8 11	15 20	Х Ү	10 10	27 27			
24058_2	16 24	14 14.2	17 17	8 11	15 20	X X	9 10	20 27			
24058_3	18 20	13 14.2	17 18	8 11	15 15	Х Ү	10 12	22 27			
Lab Code	D2S441	D22S1045	SE33	D10S1248	D1S1656	D12S391	DYS391	Yindel			
24058_1	12 14	11 15	26.2 33.2	14 15	13 14	18 20	11	1			
24058_2	12 12	15 16	28.2 33.2	13 15	14 16	19 20	-	-			
24058_3	14 14	11 17	26.2 28.2	13 14	13 15.3	18 20	11	1			

24058_3 14 14 11 17 26.2 28.2 13 Identification necessitates a comprehensive strategy that takes into account all accessible scientific and contextual information.⁵ In this instance, we employed a forensic medicine technique known as an autopsy, along with a forensic anthropology method involving visual examination of the bones and determine if they conform to human anatomy based on their size, shape, and bony characteristics. In

addition, a DNA analysis was conducted.

The field of medicine acknowledges the autopsy as an essential component.6 In addition to determining the definitive diagnosis, the autopsy establishes the relationship between the cause of death and the accompanying disorders, elucidating the interplay between the two.7 Autopsies may be classified into two primary categories: forensic autopsies and clinical autopsies. The initial kind of autopsy is conducted in instances involving suspicious, violent, or unexplained reasons of death. The pathologist conducts the second procedure in the hospital, with the permission of the deceased's closest living relative, in order to ascertain and obtain a more comprehensive comprehension of the underlying factors contributing to the individual's demise. An autopsy is a postmortem examination conducted to ascertain the origin, method, and presence of sickness or damage in a deceased corpse.8 In this instance, there was blood infiltration across the majority of the posterior region of the skull, along with evidence of severe force applied to the 4th cervical vertebra. This evidence suggests that the demise of the deceased was not a result of natural causes, but rather points to the possibility of a deliberate act of murder.

The fundamental goal of doing forensic anthropological analysis in a medicolegal setting is to establish a conclusive scientific identification of human remains. Anthropological work encompasses several tasks that aid in narrowing down the search for missing individuals. These tasks include searching for and recovering remains, identifying the species, assessing the sex, determining the age at death, measuring the stature, calculating the time since death, determining the ancestry, and identifying unique anatomical features.⁹ Our identification of the skull was successful. It belonged to a male individual of the Mongoloid race, aged between 21 and 39 years. The head and neck exhibited indications of physical aggression, indicating a time of demise exceeding 10 days. This information confirms the discovery of a decapitated corpse that occurred one week earlier.

Forensic DNA analysis refers to the methodical study and interpretation of genetic material with the purpose of identifying a perpetrator or victim, particularly in criminal investigations. Forensic DNA analysis involves the use of scientific techniques to extract, purify, and examine DNA derived from biological samples collected from suspects, victims, and crime scenes. The analysis can employ various approaches, such as DNA sequencing, polymerase chain reaction (PCR), and STR (short tandem repeat) analysis. The potential categories of specimens include blood, tissue, bones, saliva, semen, hair, and several other biological fluids.¹⁰ For this example, we obtained a DNA sample from the maxillary first premolar. Additionally, we obtained blood samples from both parents for comparison. The DNA test results concluded that X1 was the biological child of X2 and X3. Thus, science has proven that the found head bone is an integral part of a buried headless body.

CONCLUSION

Forensic medical examination, anthropology, and genetic profiling play a crucial part in both identifying victims and determining the cause of death. Autopsy and anthropological analysis can ascertain an individual's distinctive profile in this scenario. Subsequently, a genetic profile, obtained both directly and indirectly, can be employed to establish the identity of the discovered skull bones.

ETHICS APPROVAL

The Ethical approval statement should be provided including the consent. If not appropriate, authors should state: *"Not required."*

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COMPETING INTERESTS

All the authors declare that there are no conflicts of interest.

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UNDERLYING DATA

Derived data supporting the findings of this study are available from the corresponding author on request.

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