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Assessment of Gel Diffusion as a Rapid Forensic Tool for Species Identification in Processed Meat

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Abstract: Species identification in meat products is crucial for forensic investigations involving food fraud, illegal wildlife trade, and regulatory enforcement. This study evaluated the efficacy of immune diffusion (Ouchterlony technique) for identifying beef, chicken, and goat meat in raw, boiled, and cooked samples. Fresh meat specimens were procured and processed under standardized conditions. Protein extracts from raw, boiled (100°C/10 minutes), and cooked (100°C/20 minutes) samples were subjected to immunodiffusion using species-specific antisera in agarose gel plates, followed by incubation at 4°C for 72 hours.

Results demonstrated clear precipitin lines for all raw samples, confirming successful species identification through specific antigen-antibody binding. However, all thermally processed samples yielded consistently negative results across all species. This observation is attributed to irreversible heat-induced denaturation of conformational protein epitopes at 100°C, rendering proteins immunologically undetectable despite their physical presence.

The findings establish that while immunodiffusion is effective for fresh, unprocessed meat, it is unsuitable for cooked or boiled samples. The study recommends differential analytical protocols: immunodiffusion for raw meat analysis and DNA-based techniques for processed samples, ensuring accurate and legally admissible forensic determinations in food authenticity investigations.

Keywords: Species Identification, Immunodiffusion, Gel Diffusion, Forensic Science, Processed Meat, Protein Denaturation, Food Authentication

I. INTRODUCTION

Species identification constitutes a fundamental aspect of forensic biology and serology, serving as a critical tool for providing scientifically validated evidence in legal proceedings. The principle underlying species-specific protein analysis involves the separation of proteins from biological samples, such as blood or tissue, through gel electrophoresis. This technique is subsequently integrated with immunoelectrophoresis, wherein antiserum containing antibodies against a known species is allowed to diffuse from a parallel trough. When the separated serum proteins from an evidentiary sample correspond to the antibodies, precipitin arcs emerge within the gel matrix, creating a distinctive species-specific "fingerprint." This pattern definitively establishes the animal source of the sample, proving particularly valuable in cases involving poaching, illegal wildlife trafficking, and food fraud investigations.

Species identification, in its broadest scientific context, refers to the methodological process of determining an organism's biological classification at the species level through examination of diagnostic physical, genetic, or behavioral characteristics. Contemporary approaches increasingly emphasize molecular techniques, particularly DNA barcoding utilizing standardized gene regions such as mitochondrial cytochrome c oxidase subunit I (COI), which generates a genetic fingerprint unique to each species. Accurate species identification is indispensable across multiple domains, including forensic animal investigations, conservation planning, ecological research, biodiversity assessment, and monitoring of illegal trade activities. Furthermore, it facilitates the discovery of cryptic species, resolves taxonomic ambiguities, and provides essential data for understanding ecological processes and informing evidence-based management decisions.

Electrophoresis, the movement of charged molecules through an aqueous medium under an applied electric field, serves as the foundational technique for protein separation. Migration velocity is governed by both the strength of the electric field and the molecular charge, enabling differently charged molecules to form distinct zones during migration. To minimize zone diffusion and achieve high-resolution fractionation, electrophoresis is conducted in an anticonvective medium such as a gel matrix or viscous fluid, wherein molecular size also influences migration speed. Agarose, a polymer derived from red seaweed, is commonly employed as the gel medium due to its exceptional mechanical stability and tunable pore sizes, ranging from 150 nm at 1% concentration to 500 nm at 0.16% concentration, permitting effective separation of nucleic acid fragments between 400 and 23,000 base pairs.

This serum-antiserum test, commonly referred to as the precipitin test, represents an established immunological technique for determining the species origin of animal flesh.

This method exploits the specific interaction between proteins (antigens) and antibodies. Antiserum is produced by immunizing a host animal, such as a rabbit, with purified serum proteins from a known species, generating species-specific antibodies. When a protein extract from an unknown meat sample is diffused along side the antiserum in an agar gel using the Ouchterlony technique, a visible white precipitate line forms if the meat proteins match the antiserum's specificity, confirming the presence of the target species. Prior to the advent of modern DNA technologies, this immunological approach was instrumental in food authenticity verification and forensic animal investigation.

Meat species authentication holds critical importance for ensuring food safety, regulatory compliance, and consumer protection. Adulteration constitutes economic fraud while simultaneously endangering individuals with dietary restrictions or allergies, particularly when premium meats such as beef and goat are replaced with inferior alternatives. In processed products where traditional morphological identification is no longer feasible, reliable analytical techniques become imperative for species verification. Immunological methods, including serum-antiserum testing, offer a traditional yet effective solution by leveraging the exceptional specificity of antigen-antibody responses. This thesis aims to evaluate the feasibility of developing specific antisera against serum proteins of beef, poultry, and goat, and subsequently employing immunodiffusion and immunoelectrophoresis to establish a dependable and user-friendly protocol for unambiguous species identification in both raw and processed meat samples.

II. RATIONALE

Accurate biological material identification is crucial for criminal investigations, consumer protection cases, and regulatory enforcement in forensic science. The forensic toolkit for examining cooked or thermally changed meat samples—evidence frequently found in cases of food fraud, tampering, or illegal slaughter—has a major deficiency. While protein-based methods such as gel electrophoresis (SDS-PAGE) and immunological assays (antiserum tests) provide advantages for damaged or heat-processed samples where DNA may be substantially fragmented, DNA-based approaches are recommended for raw tissue.

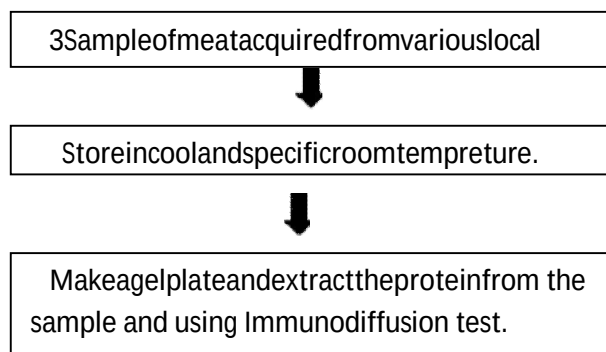
Validated comparison data on the application of these protein techniques to cooked/boiled beef, poultry, and goat meat is lacking in current forensic protocols. In real-world evidentiary circumstances, including evaluating processed food from crime scenes or confirming claims in economic fraud instances, this makes it more difficult to reliably authenticate species. It is therefore crucial to determine the sensitivity, specificity, and limits of these readily available methods for thermally treated evidence.

By methodically assessing electrophoresis and immunoassays for species identification in both raw and cooked stages, this study will directly address this forensic need. The results will improve the integrity of evidence in food-related cases, bolster forensic laboratory capabilities, and give judges more comprehensive analysis alternatives.

Some points are the research aim of our research:-

- 1) Create Standard Operating Procedures (SOPs) for handling evidence of cooked meat.
- 2) Ascertain each method's evidential threshold, or the lowest detectable amount and level of processing that still allows for identification.
- 3) To facilitate pattern matching, create reference protein/antigen profiles for important species in both raw and cooked forms.
- 4) Evaluate the methods' suitability for combination analysis, atypical situation in commercial ground beef products connected to financial fraud.

III. MATERIALS AND METHODS



A. Sample Collection

The purpose of sample collection was to get representative, fresh beef samples from typical commercial suppliers. Three different kinds of meat were obtained: goat (*Capra aegagrus hircus*), chicken (*Gallus domesticus*), and beef (*Bos taurus*). Each sample was bought from a separate local retail establishment, carefully chosen from a pool of supermarkets, specialty meat distributors, and butcher shops in the vicinity, to guarantee a varied sourcing profile and reduce batch-specific bias. To standardize the initial time-zero condition, all samples were collected on the same day.

Samples were delivered to the lab in their original, sealed retail packaging as soon as they were purchased. In order to replicate a typical consumer handling situation, they were kept at room temperature (around 25°C) for a constant, short time after arrival before being processed. All meat items were then moved to a conventional laboratory refrigerator for medium-term storage, where they were kept at 4°C for about 24 hours (overnight). The samples were stabilized throughout this storage phase, which made it possible for all experimental procedures to begin consistently.

The chilled samples were prepared under controlled, aseptic conditions before analytical or cooking processes. Each primary sample was carefully weighed into parts of around 10 grams using a sterile analytical balance. These subsamples were put into sterile, pre-labeled individual containers. During this portioning, sterile cutting boards, scalpels, and forceps were used to avoid cross-contamination between different types of meat and external microbial contamination. Each sample's integrity for further analysis was guaranteed by this cautious handling.

Laboratory-grade agarose was used to construct the gel plate on a glass slide. The Forensic Science Laboratory in Ramanagar, Varanasi, provided the particular serum and antiserum reagents. Following solidification, wells were created, and the obtained serum and antiserum were added for the ensuing electrophoresis or immuno-diffusion test.

IV. METHODOLOGY

First, make the agarose gel plate by heating distilled water to dissolve the agarose powder. Evenly pour the heated mixture onto a level glass plate, then let it harden. Agarose sets at room temperature in roughly half an hour as opposed to being refrigerated. After the gel is solid, make tiny wells in it using a "well rod" or well-cutting instrument. Fill the wells with various animal antisera, such as goat, beef, or poultry antisera. Antigens and antibodies disperse across the gel via methods such as immunodiffusion or immunoelectrophoresis.

Immune reactivity is indicated by visible precipitation lines that appear where matched pairs converge. This enables the detection of antigen-antibody interactions in a semi-solid matrix as well as the comparison of antiserum specificity.

The test's first stage entails the methodical isolation of proteins from samples of cooked and boiled beef for later immunological testing. Six sanitized China plates are prepared and arranged for this process. The first three dishes—one for chicken, one for goat meat, and one for beef—are set aside

for the cooked meats samples. Every sample of raw beef is put into the appropriated dish and labeled appropriately. After that, the samples are boiled for ten minutes at 100 degrees Celsius in distilled water. In addition to denaturing the proteins, this standardized heat treatment may mimic a particular type of food processing. The cooked meat is carefully removed after boiling, and the soluble proteins that were released into the aqueous media are gathered. As an alternative, proteins can be extracted straight from the solid tissue using instruments like forceps or a scraper, then homogenized in an appropriate solution to produce a protein extract. This procedure produces the first batch of test solutions made from unseasoned but heat-treated meat.

The cooked beef samples are then prepared using an identical set of three China dishes. Here, typical culinary ingredients like cooking oil, water, and a mixture of spices are used to cook the same kinds of meat—beef, chicken, and goat. In order to guarantee complete preparation, this cooking procedure is carried out at 100 degrees Celsius for a duration of twenty minutes. The objective is to produce a more complex, gastronomically relevant sample that contains possible chemicals from the oils and spices in addition to heat-denatured proteins. This seasoned meat's protein is removed once again after cooking. Carefully decant the liquid broth, which is full of dissolved proteins and other materials. A second, more sophisticated set of protein extracts is prepared for examination by scraping or extracting any remaining protein from the solid meat with a knife and combining it with the broth.

The immunodiffusion test is carried out once the protein extracts are ready. The anti-serum (a particular antibody solution, such as anti-beef serum) is pipetted into the central well of the previously prepared agarose gel plate, which has peripheral wells surrounding the center well. After that, the extracted protein samples—three from boiling meat and three from equivalent cooked meat—are put in a predetermined pattern into the nearby peripheral wells. After carefully placing the filled gel plate in a damp chamber to keep it from drying out, it is refrigerated for about three days.

The antigens (proteins) and antibodies permeate radially across the agarose matrix over this prolonged low-temperature incubation period. Large, insoluble immune complexes that precipitate out of solution and appear as fine white lines in the gel are formed when particular antibodies come into contact with their corresponding antigens. The impact of culinary processing, including the addition of spices, on the immunological reactivity and integrity of the meat proteins can be examined by comparing the precipitation patterns between boiled and cooked samples.

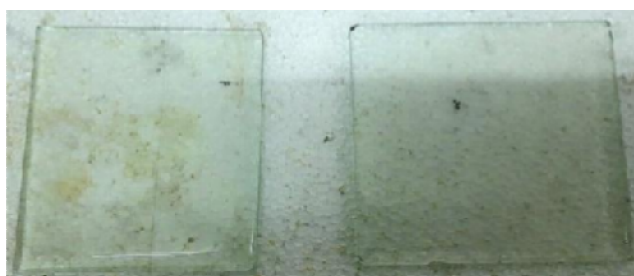
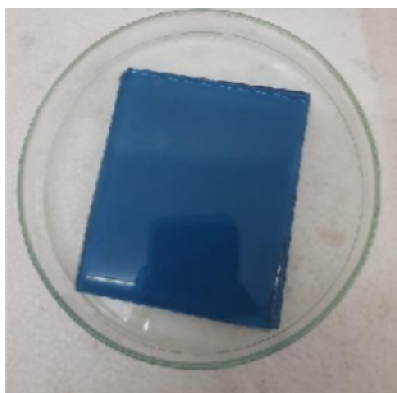


Figure1 Agarose Gel



Figure2 "Raw meat"

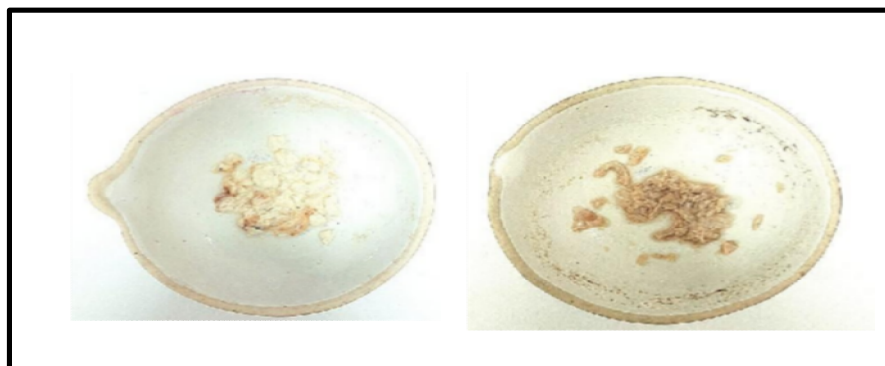


Figure3 "Boiled meat"

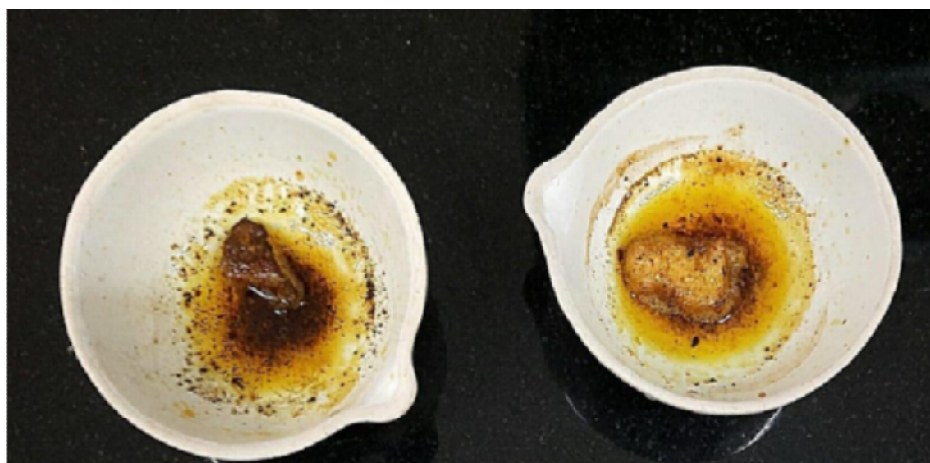


Figure4"Cookedmeat"

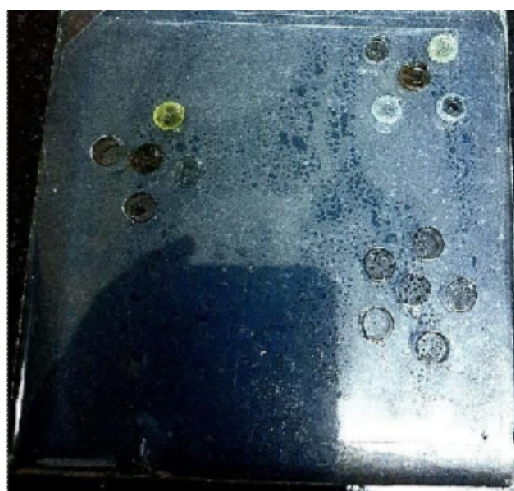


Figure5"Resultent"

ObservationTable:-

S no.	Sample	Boil	Cooked	Raw
1.	Beef	Negative(-)	Negative(-)	Positive(+)
2.	Chicken	Negative(-)	Negative(-)	Positive(+)
3.	Goat	Negative(-)	Negative(-)	Positive(+)

Result

The study used the immunodiffusion approach to distinguish between raw and boiled or cooked samples of beef, poultry, and goat meat in order to identify species. The visible precipitate that results from the binding of species-specific antibodies to their target antigens in a gel matrix is the basis for this method. The outcomes for the raw meat samples from each of the three species were consistently positive. This result was anticipated since raw beef includes a complete range of native, soluble proteins. Because these proteins maintain their exact three-dimensional structures, the antibodies in the antisera are able to identify and attach to them efficiently, creating distinct precipitate lines that verify the identity of the species.

On the other hand, all samples of cooked and boiled meat, regardless of species, produced negative results. This is a direct result of denaturation of proteins caused by heat. Meat proteins' intricate structure is irreversibly changed by the thermal processing of boiling or frying, which reveals their unique conformational epitopes. The proteins themselves remain intact, but their antibody-binding sites are either damaged or rendered unusable. As a result, the antibodies are unable to create the complexes required for precipitation. Therefore, the negative result highlights a significant shortcoming of this immunological approach for assessing processed foods: it shows an absence of immunologically reactive protein rather than an absence of protein itself.

V. RESULTS AND DISCUSSION

The immunodiffusion technique was employed to facilitate species identification through the detection of visible precipitin lines resulting from the binding of species-specific antibodies to their target antigens within the agarose gel matrix. Fresh meat samples of beef, chicken, and goat consistently yielded positive results, characterized by the formation of distinct and well-defined precipitin bands. This outcome was anticipated, as raw meat retains a comprehensive array of native, soluble proteins in their native conformational states. The preservation of these three-dimensional structures enables the antibodies present in the antisera to recognize and bind to their corresponding epitopes with high specificity and affinity, thereby generating clear precipitation patterns that conclusively verify species identity.

In marked contrast, all thermally processed samples—both boiled (100°C for 10 minutes) and cooked (100°C for 20 minutes with culinary ingredients) produced uniformly negative results irrespective of the species under investigation. This observation is directly attributable to heat-induced protein denaturation, wherein the intricate and highly ordered structures of meat proteins undergo irreversible conformational alterations. While the primary amino acid sequences remain intact, the thermal energy disrupts the secondary, tertiary, and quaternary structural arrangements, effectively destroying the conformational epitopes that antibodies recognize. Consequently, the antibody-binding sites are rendered inaccessible or structurally compromised, preventing the formation of the antigen-antibody complexes essential for precipitin formation. The negative results therefore signify the absence of immunologically reactive protein rather than the complete absence of protein itself, highlighting a significant methodological limitation of this immunological approach for the analysis of processed food products.

These findings definitively validate both the fundamental principle of immunodiffusion and its critical constraint. The technique's efficacy with raw meat confirms its dependence on the preservation of native protein antigen structures, which facilitate targeted antibody binding and visible precipitin formation. Conversely, the consistently negative outcomes for boiled and cooked samples are unequivocally attributable to heat-induced denaturation at 100°C, which irreversibly destroys the conformational epitopes recognized by the antibodies, rendering the proteins immunologically undetectable despite their physical presence.

A comparative analysis with existing literature provides valuable insight into the temperature-dependent nature of this limitation. The divergence from Tsuneo Abe's 1973 study, which reported retained antigenicity upon heating to 70°C, is particularly instructive. This discrepancy establishes that the utility of immunoassays is governed by a precise temperature threshold rather than a binary phenomenon. For the serum protein examined in the present investigation, this threshold lies between 70°C and 100°C. Consequently, this research refines the methodological application by demonstrating that immunodiffusion is unsuitable for fully cooked or boiled evidence, though it may retain applicability for minimally processed products—a distinction of paramount importance for the development of evidence-based forensic protocols.

The forensic significance of this research is substantial, as it establishes the operational limits of a conventional analytical technique, thereby preventing its inappropriate application. For the forensic analyst, a negative result from a cooked meat sample must be interpreted as an anticipated methodological failure rather than evidence of species absence. Application of this test to heat-processed evidence in cases involving food fraud, adulteration, or illegal wildlife trade could yield disastrous false negatives, potentially enabling perpetrators to evade legal accountability. This study provides the empirical foundation for excluding immunodiffusion from the analytical workflow for any heat-treated evidentiary material.

Consequently, this investigation directly supports the modernization of forensic laboratory practices by confirming the necessity of transitioning to more robust analytical methodologies for processed materials, such as DNA barcoding or polymerase chain reaction (PCR)-based techniques, which target heat-stable genetic material. The findings enable forensic practitioners to establish clear and defensible standard operating procedures: raw meat evidence may be reliably analyzed using immunodiffusion, whereas all cooked or processed evidence must be subjected to genetic analysis to ensure accurate, reliable, and legally admissible outcomes in food authenticity and wildlife crime investigations.

VI. CONCLUSION

The core concept and main drawback of immunodiffusion for meat species identification are definitely shown by this study. The technique is quite successful and produces very positive findings for fresh samples of goat, chicken, and beef. Unprocessed tissue's native proteins maintain their unique three-dimensional structures, which enable antibodies to identify and attach to them, creating distinct precipitin lines in the agarose gel. This demonstrates the technique's fundamental immunological specificity and justifies its usefulness as a trustworthy assay for species authentication in fresh, unaltered meat products, yielding a clear and unambiguous result.

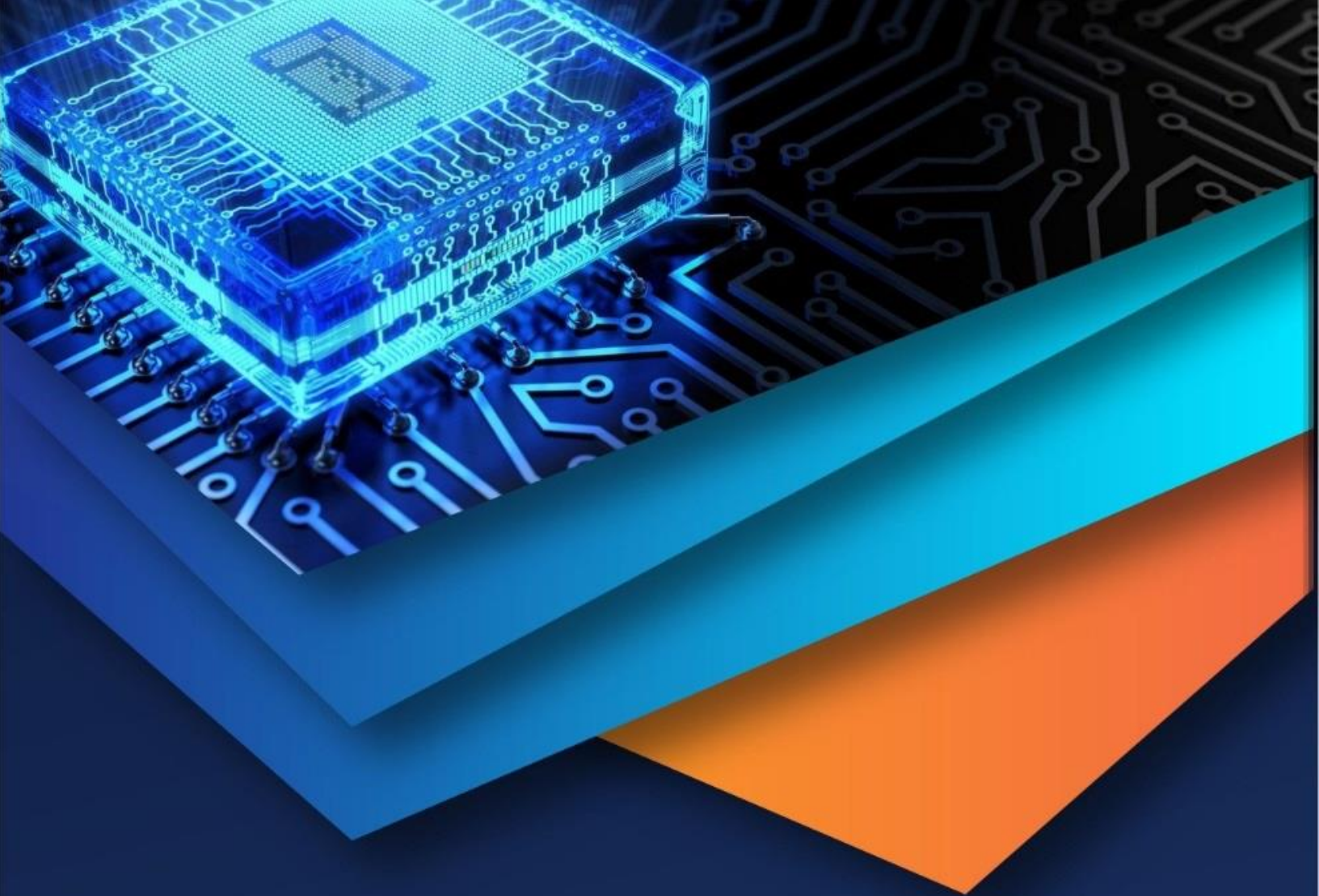
On the other hand, every sample that was heat-processed—boiled or cooked—produced unfavorable outcomes. Protein denaturation is the direct source of this important discovery. Meat proteins' conformational shape is irreversibly changed by thermal processing at 100°C, which eliminates the specific epitopes that antibodies are meant to target. Proteins become immunologically invisible even though they are physically there. Therefore, the lack of a precipitin line indicates a loss of antigenic reactivity rather than a loss of protein content, which is a false negative for species presence.

The findings are consistent with previous research, demonstrating that conventional immunological techniques are appropriate for processed meats. Heat is the primary inactivating agent, as evidenced by the unfavorable result for cooked samples, independent of additional spices or oils. The majority of forensic or food fraud evidence is cooked, tinned, or otherwise processed meats where morphology is lost, hence this poses a significant practical limitation. The method's practical forensic use is limited by its inability to bridge the gap between raw and cooked analysis.

Thus, method selection depending on sample condition is strongly recommended by this research. Laboratories must use different methods for processed evidence, even though immunodiffusion is still valid for raw meat analysis. Accurate species identification in cooked food requires the use of DNA-based techniques such as PCR, which targets heat-stable genetic material, or sophisticated proteomics, which looks for thermally resistant peptide biomarkers. This guarantees trustworthy, admissible evidence in all forensic and food authenticity inquiries.

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