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Discrimination of mongoose hair from domestic cattle hair, human hair, and synthetic fiber using FTIR spectroscopy and chemometric analysis: a rapid, cost-effective, and field-deployable tool for wildlife forensics†

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Mongoose hair is used to prepare fine brushes, which increases the demand for mongooses to be poached from the wild and brutally bludgeoned to death. Mongooses were listed as Schedule I species under the Indian Wildlife (Protection) Act 1972. Species identification of wildlife case-related samples is necessary to convict a person under this legislation. Microscopy and DNA-based techniques are commonly used to identify mongoose hair in seized brushes. However, in painting brushes, the roots, and the lower part of the hair are mostly trimmed, and only the upper part is used to make the brushes. In addition, brushes are often prepared with mixed hair from mongoose, domestic cattle, human hair, and synthetic fibre. Therefore, the identification of mongoose hair by microscopy and DNA-based techniques is restricted due to the lack of complete strands of hair and the absence of hair roots. Therefore, there is an urgent need to develop an alternative methodology for the identification of mongoose hair from seized articles. FTIR spectroscopy for forensic analysis has gained significant attention over the years because of its sensitivity, specificity, and non-destructive nature. The present study aimed to discriminate Indian grey mongoose (*Herpestes edwardsii*) hair from domestic cattle hair (domestic water buffalo and domestic cow), human hair, and synthetic fiber based on their chemical composition using FTIR spectroscopy and chemometric analysis. We have taken hair from four individuals for each species, namely Indian grey mongoose, domestic cattle, human hair, and synthetic fibre. The FTIR spectrum was recorded, and partial least-squares discriminant analysis (PLS-DA) was used to discriminate hair and synthetic fiber. The established PLS-DA model showed an R-square value and an RMSE (root mean square error) value of 0.9 and 0.13 respectively. Our preliminary findings have shown that FTIR spectroscopy combined with chemometrics can quickly discriminate Indian grey mongoose hair, domestic cattle hair, human hair, and synthetic fiber, providing crucial evidence for judicial proceedings.

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1. Introduction

India is a megadiverse country, home to 7–8% of the world's flora and fauna. It accommodates more than 45 000 species of plants and 91 000 species of animals.¹ Wildlife in India is protected by the Wildlife (Protection) Act 1972 (WPA), which prohibits the trade of more than 1800 species of wild animals,

plants, and their derivatives.^{2–4} Despite strong legal protection, poaching and illegal trade have been increasing over the years, posing a significant threat to several iconic species and pushing them to the brink of extinction. Mammals are frequently poached for their skin, meat, hair, and wool, which are widely traded for different purposes, such as fur coats, leather purses, belts, shoes, carvings, *etc.*^{5,6}

In recent years, mongooses (*Herpestes* sp.) have been poached illegally and their hair is used to produce painting and shaving brushes.^{7,8} The various seizures made by law enforcement agencies indicate that poaching of these animals is still regular and affects the conservation of these animals.⁷ For example, in 2023, the Andhra Pradesh Forest Department seized more than 18 000 paint brushes from several premises allegedly used by dealers in Hyderabad city, and forensic analysis confirmed that the paint brushes were prepared with Indian

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grey mongoose (*Herpestes edwardsii*);⁹ The Tamil Nadu Wildlife Crime Control Bureau seized 14 000 mongoose hair paint brushes from various districts of Tamil Nadu, such as Chennai, Coimbatore, and Madurai.¹⁰ The Wildlife Crime Control Bureau (WCCB), India, has conducted enforcement operations in seven Indian states from 2002 to 2021, leading to the recovery of more than 1.6 lakh mongoose hair paint brushes and 292.7 kg of raw mongoose hair.¹¹ Additionally, the Centre for Wildlife Forensic Sciences, Advanced Institute for Wildlife Conservation, Tamil Nadu Forest Department, Vandalur, Chennai, has received several cases involving the use of mongoose hair. A study documented that there are seven mongoose species in India⁷ namely (1) *Herpestes edwardsii*, (2) *Herpestes smithii*, (3) *Herpestes urva*, (4) *Herpestes palustris*, (5) *Herpestes javanicus*, (6) *Herpestes vitticollis* and (7) *Herpestes brachyurus*. All mongoose species are protected under the Indian Wildlife (Protection) Act 1972 and classified as Schedule I species,¹² and the International Union Conservation of Nature (IUCN)¹³ categorized mongoose species as 'Least Concern'.

In general, microscopy was used to identify mongoose hair in the seized brushes. The Wildlife Institute of India (WII)¹⁴ has demonstrated the identification of Indian grey mongoose from guard hair. The Zoological Survey of India (ZSI)¹⁵ and the Wildlife Crime Control Bureau (WCCB)¹⁶ have reported microscopic structures of dorsal guard hairs for identification of mongoose species. However, the ZSI report lacks detailed descriptions of microscopic hair characters, and the WCCB report has no microscopic images. As a result, the practical application of these findings for forensic case analysis is limited. To address this issue, Sahajpal *et al.* (2009)⁷ demonstrated the identification of four mongoose species (*H. edwardsii*, *H. smithii*, *H. palustris*, and *H. urva*) by microscopic examination of the pattern of the dorsal guard hair band and discriminate functional analysis (DFA). However, all studies have exhibited the identification of mongoose species by microscopy examination of the dorsal guard hair band pattern. Still, hairs from other regions of the mongoose body (*e.g.*, tail) are also used in brush manufacturing. Therefore, further studies must be employed and compared with the reference data to overcome this limitation. Moreover, the identification of species from hair is subject to several factors, such as the availability of complete strands of hair, the expertise of the forensic examiner, the quality and quantity of the hair, the availability of known samples (reference hair samples) and the instrumentation used for analysis.^{17–21} The results of microscopy analysis may sometimes not be conclusive; therefore, further study is needed to identify the species.²¹

Next, DNA analysis is the most common method used to detect hair at the molecular level.²² Before analysis, a forensic analyst classifies the hair based on the morphology of the hair roots to apply the appropriate analytical methods.^{23,24} In forensic analysis, hair samples with anagen follicular tissue are ideal for DNA analysis.²⁵ However, up to 95% of hair collected as forensic evidence has been documented to be telogen, which is completely keratinized and contains a small amount of cellular material.^{26,27} To overcome this issue, Bourguignon *et al.* (2008)²⁸ classified telogen hairs found at the crime scene such as type 1,

without visible soft tissue; type 2, with a small amount of soft tissue attached; and type 3, with a large amount of soft tissue attached.

In most cases, the seized brushes received for forensic analysis were type 1 and DNA extraction; subsequent analysis is very challenging.^{7,22} However, few studies have reported the identification of species on hair shafts, though the success rate is subject to various parameters, including the content of keratin.²⁹ In addition to this, during our forensic examination of the brushes received for analysis, we observed that the brushes prepared with mongoose hair were often mixed with hair from other domestic cattle, such as domestic water buffalo (*Bubalus bubalis*), domestic cow (*Bos taurus*), human hair, and synthetic fibers. Therefore, the identification of mongoose hair from seized brushes is a very tough task using microscopy and DNA-based techniques due to three primary reasons: (1) lack of complete hair structure (hair strands), (2) the hair samples received for analysis are telogen, and (3) the seized brushes often mixed with hair from domestic cattle, human hair, and synthetic fibres. Therefore, it is a fact that identifying mongoose hair from seized brushes and discriminating it from counterfeit (*i.e.*, hair from domestic cattle, humans, and synthetic fibres) is a major concern, and there is an urgent need to develop robust techniques/methods to distinguish mongoose hair from counterfeit materials.

The application of vibrational spectroscopy for forensic purposes has gained significant attention over the years because of its sensitivity, specificity, and non-destructive nature.³⁰ The infrared spectrum displays the vibrational characteristics of a sample based on the different absorption frequencies of the individual functional groups present in the sample (*e.g.*, based on the chemical composition of the sample).³⁰ In light of this, FTIR is used to analyse various materials in forensic science, including biomedical samples,³¹ paint,³² fingerprints,^{33,34} and ink.³⁵ Furthermore, studies have documented that the identification of species from hair using Attenuated Total Reflectance Fourier Transform Infrared Spectroscopy (ATR-FTIR).^{36–40} Notably, Manheim *et al.* (2015)⁴¹ described the identification of human, animal, and synthetic fiber by ATR-FTIR spectroscopy. Boll and co-workers (2017)⁴² researched the use of ATR-FTIR spectroscopy for the classification of dyed and non-dyed hairs; Bhatia *et al.* (2024)⁴³ showed the identification of hairs from Royal Bengal Tiger (*Panthera tigris tigris*), Indian Leopard (*Panthera pardus fusca*) and snow Leopard (*Panthera uncia*) using the ATR-FTIR spectroscopic technique in combination with chemometric analysis. Hence, previous studies have well documented that species, biological, and non-biological articles can be identified by ATR-FTIR spectroscopy and chemometric analysis. Therefore, we have been motivated to employ FTIR spectroscopy combined with chemometric tools for the identification of mongoose hair from counterfeit hair (*i.e.*, hair from domestic cattle and human).

In light of this, the current study demonstrates FTIR spectroscopy combined with chemometric tools for the identification of Indian grey mongoose hair (*Herpestes edwardsii*) and discriminating it from domestic cattle, such as domestic water buffalo (*Bubalus bubalis*), domestic cow (*Bos taurus*), human



hair, and synthetic fiber. We selected the Indian grey mongoose for this study because it is the most common mongoose found in southern India and its proximity to human habitation.⁹ As a result, Indian grey mongooses are often poached and their hair is used for brush preparation. The results of this study will play an important role in the wildlife forensics in identifying Indian grey mongoose hair from seized articles.

2. Materials and methods

2.1 Collection of Indian grey mongoose hair (voucher sample)

We collected 100 hair samples (25 hair strands from each individual) from four Indian grey mongooses in our Morphometry Laboratory at Advanced Institute for Wildlife Conservation (AIWC), Chennai, Tamil Nadu, India. No sample was collected from live animals.

2.2 Collection of domestic cattle hair samples

From the Kolapakkam Farm house, Chennai, India, approximately 25–30 tail hairs were collected from each of the four individuals of domestic water buffalo (*Bubalus bubalis*) and domestic cow (*Bos taurus*). Before FTIR spectroscopy analysis, the species was confirmed by molecular analysis.

2.3 Human hair collection

Human hair samples were obtained for this study from four volunteers willing to donate a small amount of scalp hair. For all hair donors, only their age and biological sex were recorded; donor names and all other personally identifiable information were not collected. The donated hairs were visibly found to be natural (non-dyed) and black.

2.4 Synthetic fiber

In this study, we used polyethylene terephthalate fiber. For FTIR analysis, we chopped small pieces of fiber and prepared the samples as described below.

2.5 Sample preparation for FTIR spectroscopy analysis

About 25 hair strands were taken from each individual, chopped into small fragments and finely powdered using a sterile mortar and pestle. The powdered samples were then thoroughly mixed with 1.0 g of KBr powder (Merck Millipore, IR spectroscopy grade) and a 13 mm KBr pellet was prepared. The mortar and pestle were cleaned with acetone (analytical grade) and then dried before and after the preparation of the sample to avoid cross-contamination.

2.5.1 Instrumentation and spectral collection. FTIR spectra were recorded using Bruker FTIR model: Alpha II (Bruker Optics, Ettlingen, Germany) equipped with a ZnSe crystal accessory and a DTGS detector. A blank KBr pellet without the sample was run for the background scan before switching to the subsequent sample. The sample holder and instrument accessories to prepare the pellet were cleaned with analytical grade acetone after each experiment to avoid cross-contamination. The spectra

were measured in transmittance mode within the range 4000–500 cm^{-1} (mid-infrared region) with a spectral resolution of 4 cm^{-1} . All experiments were averaged from 24 scans and performed at room temperature. Each sample was analysed in triplicate and a total of 48 reference spectra were recorded (16 samples $\times$ 3 = 48). Furthermore, we also recorded the FTIR spectrum for four synthetic fibers (4 samples $\times$ 3 = 12). In total, we recorded 60 spectra for this study.

2.6 Data analysis

2.6.1 Spectra preparation, analysis, and comparison of spectra. The FTIR spectral data obtained from the OPUS software (Version 8.8, Bruker Optics, Ettlingen, Germany) were pre-processed within the software. Atmospheric compensation was applied to reduce the influence of the CO_2 and H_2O absorption bands, and a baseline correction was performed on every spectrum before any analysis. Graphical plots for the reference and suspected samples were created using Origin Pro 9. Then, the functional groups present in the primary compounds in the hair were annotated.

2.6.2 Chemometric analysis. Chemometrics analysis is an interdisciplinary field of applied mathematics and statistics that helps to design, gather, analyse, and validate the information from an intrinsic set of data obtained from various analytical techniques.⁴⁴ In our study, the partial least squares discriminant analysis (PLS-DA) is used to analyse the spectral data. PLS-DA is a linear classification method that combines the properties of partial least squares regression and the discriminative ability of classification technology.⁴⁵ Hence, the current study used the PLS-DA model to discriminate Indian grey mongoose hair from human hair, synthetic hair, domestic cattle. Before applying the PLS-DA model, the data was pre-processed using the Savitzky–Golay algorithm for smoothing, the Mean centering method, and normalisation. A classification model was constructed using PLS-DA to classify Indian grey mongoose hair, domestic cattle, human hair, and synthetic fiber. Four samples were taken from each species to construct the calibration set of the PLS-DA model. All data were analysed using MATLAB R2020b (The MathWorks, MA, USA) equipped with PLS Toolbox 9.3 (Eigenvector Research, WA, USA).

3. Results and discussion

3.1 FTIR spectroscopy analysis for Indian grey mongoose hair, domestic cattle hair, human hair, and synthetic fiber

Identifying seized wild animal parts is a great challenge due to the many species involved in the illegal wildlife trade and the various forms of seized items. These articles are often made from materials derived from domestic animals and other sources (or materials); therefore, the species identification more complex using conventional methods.⁴⁶ For example, CITES has documented that wild mammal skins (leather, fur) and products (hair, wool) are widely traded, often mixed with counterfeit furs, making the identification of wildlife products challenging.⁴⁷ Therefore, there is an urgent need to develop alternative methods to identify intra- and interspecies variations in



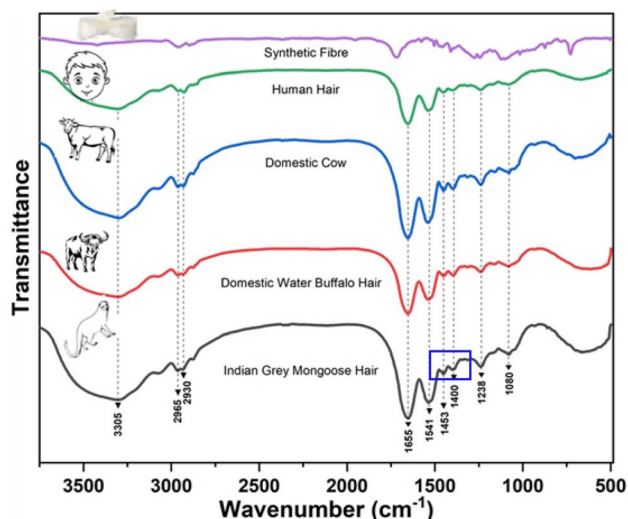


Fig. 1 FTIR spectra of hair from Indian grey mongoose, domestic water buffalo, domestic cow, human hair and synthetic fiber. The inserted clipart's were collected from online sources (for better visualization purposes), and Microsoft PowerPoint was used to prepare the figure. The blue box indicates the spectral differences in mongoose hair compared to other animal hair used in this study (including human hair).

wild and domestic species. This will enable a better implementation of national and international laws to effectively control wildlife trade.

In this study, FTIR spectroscopy was used to discriminate Indian grey mongoose hair, domestic cattle hair, human hair, and synthetic fiber. Hence, we recorded the FTIR spectrum for the hair of the Indian grey mongoose, domestic cattle hairs, human hair and synthetic fibres. The findings revealed that we could visually distinguish synthetic fiber from hair (all the hair used in this study). The results suggested that through FTIR analysis, visually can be confirm seized brushes contain hair or any other materials (including other than hair origin). On another hand, a slight spectral difference, especially in the amide-II region, was observed in the hair of Indian grey mongooses compared to domestic animals (Fig. 1). Similar results were observed in the FTIR spectrum recorded for intra-species (different individuals from the same species)

(Supplementary Fig. S1†). The primary biochemical component of animal fibres is keratin, a fibrous structural protein widely found in human and animal organs, including the epidermis, hoof, horn, hairs, and feathers.⁴⁸ Gallagher (2009)⁴⁹ detailed the protein composition of some animal fibers, while Espinoza *et al.* (2009),⁵⁰ Manheim *et al.* (2016),⁴¹ Agarwal *et al.* (2019),⁵¹ Wang *et al.* (2021),⁴⁸ and Xu *et al.* (2022)⁵² characterized hair proteins using spectroscopy techniques and described their chemical composition and corresponding wavenumbers (Table 1).

In our study, the peak observed at 3305 cm^{-1} is related to the organic material amide A (primary amide) from the overlap of the N-H and O-H hydrogen amide regions (NH stretching vibrations). Based on previous studies on keratin, the peaks at 2965 cm^{-1} and 2930 cm^{-1} represent asymmetric and symmetric $\nu(\text{CH}_3)$, respectively, and amide I is identified at 1655 cm^{-1} which corresponds to the stretching vibrations of the C=O bond (amide group). The detected peak at 1541 cm^{-1} is the amide II band that arises from vibrations of NH bending and C-N stretching, and the narrow peaks at 1453 cm^{-1} and 1400 cm^{-1} are due to the deformation in $\delta(\text{CH}_2)$ (CH_3) and $\delta(\text{CH}_3)$ respectively. Similarly, a weak peak at 1238 cm^{-1} illustrates amide III, a combination of stretching vibrations of the C-N and C-O bonds and bending vibrations of the N-H and O=C-N bonds. As reviewed and documented by Espinoza *et al.*, 2009, the region between 1200 to 1000 cm^{-1} is associated with vibrations of the sulfur-oxygen groups of keratins, and hence the peak at 1080 cm^{-1} is of $\nu_s(\text{SO})$ from cysteine oxide.⁵⁰ Based on the FTIR spectra recorded for hair samples, we found no visual difference in the spectra (except Indian grey mongoose hair), as all hair is made of keratin proteins. Additionally, relying on visible observation of FTIR spectra can lead to observer bias. To address these issues, we used chemometric analysis to distinguish Indian grey mongoose hair from domestic cattle hair, human hair, and synthetic fiber.

3.2 Chemometric modelling for discrimination of Indian grey mongoose hair, domestic cattle hair, human hair, and synthetic fiber

The PLS-discriminant analysis (PLS-DA) method is commonly used for predictive, descriptive modelling, and discriminative variable selection.⁵⁴ PLS-DA has various applications in various

Table 1 Assignment of IR absorption bands in the spectra of hair

S. No.	Wavelength (cm^{-1})	Assignment	Source	Ref.
1	3100–3500	Amide A (overlap of amide hydrogen N-H region and O-H region)	N-H stretching vibrations	48–53
2	2850–3000	$\nu(\text{CH}_3)$ asymmetric; $\nu(\text{CH}_3)$ symmetric	Saturated and unsaturated C=H, C-H region	
4	1600–1700	Amide I (carbonyl oxygen C=O region)	C=O stretching vibrations	
5	1500–1572	Amide II (amide hydrogen N-H bond)	N-H formation vibrations and C-N stretching vibrations	
6	1361–1470	Amide III $\delta(\text{CH}_2)$ (CH_3) deformation	C-N region	
7	1210–1290	Amide III (carbonyl oxygen C-O region)	C-N and C-O stretching vibrations, N-H and O=C-N bending vibrations	
8	998–1100	Sulphur oxygen (S-O vibration, C-H bond)	Cystine oxides	



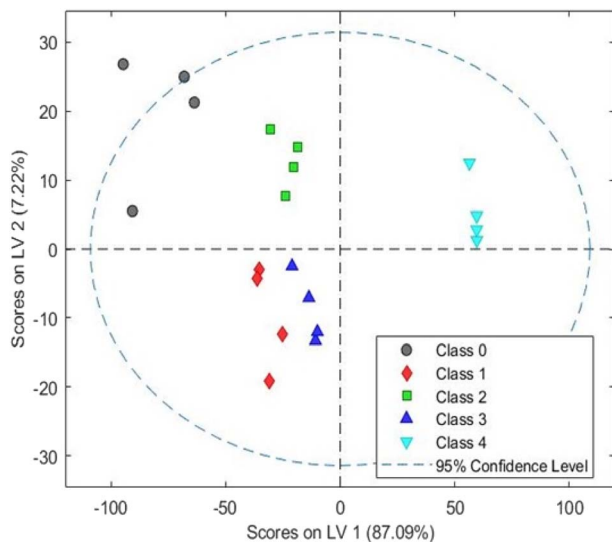


Fig. 2 Score plot for the PLS-DA classification model with LV1 vs. LV2. Class 0 – Indian grey mongoose, class 1 – domestic water buffalo, class 2 – domestic cow, class 3 – human hair and class 4 – synthetic fiber.

fields, such as forensic science, banking, medical diagnosis, food analysis, metabolomics, and soil science.^{54–58} Keeping in view, this study used PLS-DA analysis to differentiate between Indian grey mongoose hair, domestic cattle hair, human hair, and synthetic fiber based on their intensity of IR radiation absorption by biochemical composition of each hair sample.

The PLS-DA model was calculated as three latent variables (LV), namely LV1 (87.09%), LV2 (7.22%), and LV3 (2.82%); consequently, it contributed to the separation of the five groups. Score plots between LVs are shown in Fig. 2 and 3. The plot clearly showed species wise cluster and differentiation; the

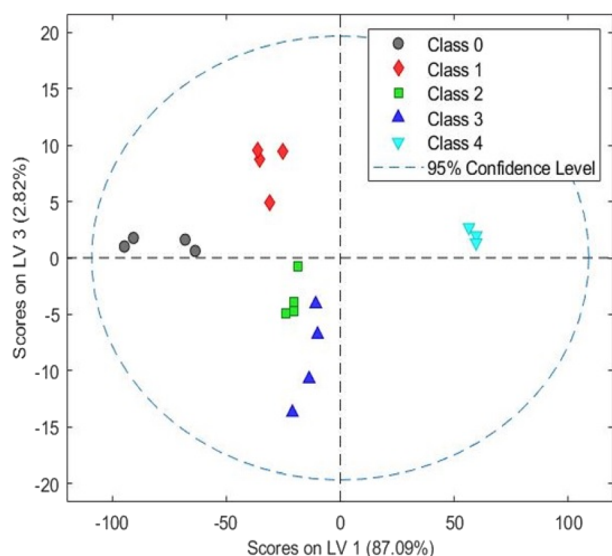


Fig. 3 PLS-DA classification model: score plot LV1 vs. LV3. Class 0 – Indian grey mongoose, class 1 – domestic water buffalo, class 2 – domestic cow, class 3 – human hair and class 4 – synthetic fiber.

latent variables effectively highlighted the differences in the FTIR spectra recorded for each hair sample. The developed model has shown an R -square value of 0.9 and an RMSE value (root mean square error) of 0.13. Hence, the PLS-DA analysis has successfully discriminated the Indian grey mongoose hair, domestic cattle hair, human hair, and synthetic fiber based on the IR transmittance variations of the chemical compositions present in the hair samples (especially keratin protein).

Proteins are made of hundreds of amino acids folded into a well-defined structure that is stabilized by thousands of interactions. Infrared (IR) spectroscopy can reveal the structure of proteins *via* vibrational resonances from the polypeptide backbone and/or side chains. Such resonances depend on the protein structure and local interactions (such as hydrogen bonding between amino acids).⁵⁹ In light of this, studies documented that FTIR spectroscopy analysis of polypeptides and proteins reveals nine characteristic IR absorption bands: amide A, B, and I–VII. The amide I and II bands are the most prominent vibrational bands of the protein backbone. The amide I band ($1700\text{--}1600\text{ cm}^{-1}$) is the most sensitive spectral region to the protein's secondary structural components and is primarily due to the $\text{C}=\text{O}$ stretch vibrations of the peptide linkages. On the other hand, the amide II band is mainly derived from in-plane NH bending and the CN stretching vibration.^{48–53}

In this regard, the current study aimed to discriminate the Indian grey mongoose hair, domestic cattle hairs, human hair, and synthetic fibres based on their biochemical composition using FTIR analysis. The FTIR analysis of hair showed the characteristic peaks at 1655 cm^{-1} (amide I), 1541 cm^{-1} (amide II), and peaks at 1453 cm^{-1} and 1400 cm^{-1} are due to deformation in $\delta(\text{CH}_2)$ (CH_3) and $\delta(\text{CH}_3)$ respectively.⁶⁰ These spectral peaks were attributed to hair keratin protein (abundant in the outer layer of skin, hair, and nails) and similar peaks were observed in all the hair samples, excluding slight differences in Indian grey mongoose hair. Studies reported that in the protein analysis through IR spectroscopy, the amide I band is sensitive to the secondary structure and is not strongly influenced by side chains of proteins. In contrast, the IR absorption properties vary in the α -helices, β -sheets, random coils, and loops (*i.e.*, protein secondary structure elements) due to hydrogen bonding of different strength from one peptide $\text{C}=\text{O}$ group to H–N– group of a neighboring amino acid. The strength of the hydrogen bond plays a vital role in the absorption frequency of the $\text{C}=\text{O}$ vibration in the amide I, corresponding to different secondary structure segments within the protein.^{61–63}

Considering that keratin protein is present in all the hair used in this study, the amino acid compositions of hairs vary from species to species. Subsequently, the hydrogen bonding patterns change within keratin proteins (keratin in different hair types). As a result, it impacted in IR absorption properties of keratin proteins present in each hair type. The small variations in amide I and amide II regions of hair cannot be identified by the naked eye. However, PLS-DA analysis identified the difference in IR absorption properties in the amide I, amide II, and amide III regions of hair and discriminated.⁶³ At this movement, the current study demonstrated discrimination of



mongoose hair, domestic cattle, human hair, and synthetic fiber using FTIR spectroscopy combined with chemometric analysis.

However, the number of samples was limited to only four in the current study; therefore, we were unable to validate the results. The study showed that FTIR spectroscopy combined with chemometric analysis could discriminate Indian grey mongoose hair, domestic cattle (domestic water buffalo and domestic cow), synthetic fiber, and human hair. The results indicate that this method can be used to identify Indian grey mongoose hair from seized articles and can discriminate it from domestic cattle hair, human hair, and synthetic fiber.

Furthermore, it is believed that FTIR spectroscopy combined with chemometric analysis will facilitate the rapid identification of Indian grey mongoose from seized articles and will be cost-effective compared to molecular techniques. Furthermore, creating an FTIR spectral library for Indian grey mongoose hair could be beneficial, as it could be utilized in the field, and border force and law enforcement officials can use it to identify mongoose hair in seized articles. This study showed positive signs for the identification of mongoose hair in seized articles, including mixed mongoose hair from domestic cattle, human hair, and synthetic fiber. However, to enhance the forensic applications of this methodology, additional studies should be performed to validate it with a large number of hair samples, which will facilitate the construction of a comprehensive classification and discriminant model.

Our study recorded the FTIR spectra of samples made with potassium bromide (KBr) pellets. As a result, we used a fine powder of 25 hair samples to record the spectra of one sample (in triplicate). Due to the non-availability of hair samples, we are unable to use a high volume of samples for this study. Consequently, the data generated are inadequate to validate this methodology. Furthermore, India is home to seven species of mongoose; our study specifically focused on identifying Indian grey mongoose hair. Therefore, future studies should aim to identify and differentiate the other mongoose species. Despite these limitations, our proof-of-concept study marks a significant advance in species discrimination based on hair analysis. The results obtained from this study will have practical applications in wildlife forensic investigations.

4. Conclusions

In wildlife forensics, discriminating or identifying individual species from an unknown sample is necessary to convict a person under the law. The current study attempted to identify Indian grey mongoose hair and distinguish it from domestic cattle hair, such as domestic water buffalo, domestic cow, human hair, and synthetic fiber, using FTIR spectroscopy combined with chemometric analysis. The preliminary results of this study documented that FTIR spectroscopy, combined with chemometric analysis, is a powerful tool for discriminating Indian grey mongoose hair from other animal hair and has great potential for forensic applications. However, the current study is carried out with a minimum sample size; as a result, we were unable to validate the results. Therefore, to apply these

methods to forensic practice, we will improve the experimental setup with a high volume of samples in the following research to resolve the problem stated above.

Ethical approval

The Principal Chief Conservator of Forests & Chief Wildlife Warden, Tamil Nadu Forest Department, Government of Tamil Nadu, has permitted to use wild animal hair samples for this study. Human hair samples were collected from volunteers after the project scientist explained the purpose of the hair collection and the importance of the project. Domestic cattle hair was collected from the farm in the presence of the Institute's veterinary doctor, and the project scientist explained the purpose of the hair collection to the farm owner.

Data availability

All of the data generated in this study is presented in this manuscript.

Author contributions

SAJ: conducted all the experiments, analyzed the FTIR data, and prepared the graphs. CB: chemometric analysis. RS: assisted in discussions and edited the manuscript. KBT: developed the idea and research methodology, analyzed the data, drafted the original manuscript and supervised. DV: logistic support. AU: funding arrangements. All authors have read and approved the manuscript for publication.

Conflicts of interest

The authors declare that they have no conflicts of interest. The authors are solely responsible for the content and writing of this manuscript.

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References

- 1 *Biodiversity and Conservation*, <https://ebooks.inflibnet.ac.in/esp03/chapter/current-status-of-biodiversity-in-india/>, accessed 20 May 2024.
- 2 L. Wilson-Wilde, *Forensic Sci. Med. Pathol.*, 2010, **6**, 221–222.
- 3 M. Anagnostou and B. Doberstein, *Ambio*, 2022, **51**, 1615–1631.
- 4 M. GoI, *Facc Minist. Environ.*, 1972, 1–218.



- 5 A. K. Rana and N. Kumar, *Biodivers. Conserv.*, 2023, **32**, 1473–1491.
- 6 A. Mozer and S. Prost, *Forensic Sci. Int. Anim. Environ.*, 2023, **3**, 100064.
- 7 V. Sahajpal, S. P. Goyal, R. Raza and R. Jayapal, *Sci. Justice*, 2009, **49**, 205–209.
- 8 Wildlife Trust of India, <https://www.wti.org.in/news/trade-in-mongoose-hair-illegal-from-today>, accessed 20 May 2024.
- 9 Wildlife Trust of India, <https://www.wti.org.in/news/huge-seizures-of-mongoose-hair-brushes-in-hyderabad>, accessed 20 May 2024.
- 10 *The Hindu*, <https://www.thehindu.com/news/national/tamil-nadu/tamil-nadu-wildlife-crime-control-bureau-seizes-14000-paint-brushes-made-of-mongoose-hair/article67238995.ece>, accessed 20 May 2024.
- 11 Wildlife Trust of India, <https://www.wti.org.in/news/massacre-of-the-mongoose-operation-to-dismantle-trade-in-mongoose-hair-brushes>, accessed 20 May 2024.
- 12 India Code, <https://www.indiacode.nic.in/bitstream/123456789/1726/1/a1972-53.pdf>, accessed 20 May 2024.
- 13 IUCN Red List of Threatened Species, <https://www.iucnredlist.org/search?query=mongoose&searchType=species>, accessed 20 May 2024.
- 14 A. Bahuguna, V. Sahajpal, S. P. Goyal, S. K. Mukherjee and V. Thakur, *Species Identification from Guard Hair of Selected Indian Mammals*, Wildlife Institute of India, 2010.
- 15 M. Kamalakannan, C. Venkatraman and K. Chandra, Trichotaxonomy of indian mammals: carnivores, artiodactyls, primates, giant squirrels, lagomorphs and pangolins, *Zool. Survey India*, 2019, 1–227.
- 16 Wildlife Crime Control Bureau, *A Manual on Wildlife Species in Trade*, 2011, <http://wccb.gov.in/WriteReadData/userfiles/file/AManualonWildlifeSpeciesintrade.pdf>, accessed 21 May 2024.
- 17 J. K. De, S. Chakraborty and R. Chakraborty, *Mammalia*, 1998, **62**, 285–296.
- 18 A. Linacre and S. S. Tobe, *Investig. Genet.*, 2011, **2**, 1–9.
- 19 M. S. Dahiya and S. K. Yadav, *J. Forensic Res.*, 2013, **4**, 178.
- 20 L. A. Hausman, *Sci. Mon.*, 1930, **30**, 258–277.
- 21 Z. Zafarina and S. Panneerchelvam, *Malays. J. Med. Sci. MJMS*, 2009, **16**, 35.
- 22 V. Sahajpal and S. P. Goyal, *DNA Fingerprinting: Advancements and Future Endeavors*, 2018, 89–127.
- 23 Z. Liu, H. Simayijiang, Q. Wang, J. Yang, H. Sun, R. Wu and J. Yan, *Int. J. Legal Med.*, 2023, **137**, 613–633.
- 24 S. L. Koch, S. R. Tridico, B. A. Bernard, M. D. Shriver and N. G. Jablonski, *Am. J. Hum. Biol.*, 2020, **32**, e23316.
- 25 E. M. Brooks, M. Cullen, T. Szytdna and S. J. Walsh, *Aust. J. Forensic Sci.*, 2010, **42**, 115–122.
- 26 J. Edson, E. M. Brooks, C. McLaren, J. Robertson, D. McNevin, A. Cooper and J. J. Austin, *Forensic Sci. Int. Genet.*, 2013, **7**, 180–188.
- 27 C. A. Linch, *J. Forensic Sci.*, 2009, **54**, 346–349.
- 28 L. Bourguignon, B. Hoste, T. Boonen, K. Vits and F. Hubrecht, *Forensic Sci. Int. Genet.*, 2008, **3**, 27–31.
- 29 K. S. Grisedale, G. M. Murphy, H. Brown, M. R. Wilson and S. K. Sinha, *Int. J. Legal Med.*, 2018, **132**, 107–115.
- 30 J. Manheim, Honors thesis, University at Albany, 2015.
- 31 S. G. Kazarian and K. L. A. Chan, *Biochim. Biophys. Acta, Biomembranes*, 2006, **1758**, 858–867.
- 32 C. Muehlethaler, G. Massonnet and P. Esseiva, *Forensic Sci. Int.*, 2011, **209**, 173–182.
- 33 M. Szafarska, M. Woźniakiewicz, M. Pilch, J. Zięba-Palus and P. Kościelniak, *J. Mol. Struct.*, 2009, **924**, 504–513.
- 34 N. J. Crane, E. G. Bartick, R. S. Perlman and S. Huffman, *J. Forensic Sci.*, 2007, **52**, 48–53.
- 35 W. Dirwono, J. S. Park, M. R. Agustin-Camacho, J. Kim, H.-M. Park, Y. Lee and K.-B. Lee, *Forensic Sci. Int.*, 2010, **199**, 6–8.
- 36 K. Virkler and I. K. Lednev, *Anal. Chem.*, 2009, **81**, 7773–7777.
- 37 H. Panayiotou, PhD thesis, Queensland University of Technology, 2004.
- 38 S. Brandes, PhD thesis, Queensland University of Technology, 2009.
- 39 P. M. J. Barton, PhD thesis, Queensland University of Technology, 2011.
- 40 D. Kurouski and R. P. Van Duyne, *Anal. Chem.*, 2015, **87**, 2901–2906.
- 41 J. Manheim, K. C. Doty, G. McLaughlin and I. K. Lednev, *Appl. Spectrosc.*, 2016, **70**, 1109–1117.
- 42 M. S. Boll, K. C. Doty, R. Wickenheiser and I. K. Lednev, *Forensic Chem.*, 2017, **6**, 1–9.
- 43 D. Bhatia, C. P. Sharma, S. Sharma and R. Singh, *Sci. Justice*, 2024, **64**, 314–321.
- 44 L. Mariey, J. P. Signolle, C. Amiel and J. Travert, *Vib. Spectrosc.*, 2001, **26**, 151–159.
- 45 D. Ballabio and V. Consonni, *Anal. Methods*, 2013, **5**, 3790–3798.
- 46 V. Sharma, C. P. Sharma, V. P. Kumar and S. P. Goyal, *Forensic Sci. Int.*, 2016, **266**, 226–233.
- 47 Using the CITES Appendices – UNIA, <https://cites.unia.es/cites/file.php/1/trainers/E-GCBriefcase-05.ppt>, accessed 20 May 2024.
- 48 X. Wang, Z. Shi, Q. Zhao and Y. Yun, *Materials*, 2021, **14**, 1–15.
- 49 W. Gallagher, W. Xu, J. Xia, S. Min and Y. Xiong, *Spectrochim. Acta, Part A*, 2022, **274**, 121034.
- 50 E. O. Espinoza, B. W. Baker, T. D. Moores and D. Voin, *Endanger. Species Res.*, 2009, **9**, 239–246.
- 51 V. Agarwal, A. G. Panicker, S. Indrakumar and K. Chatterjee, *Int. J. Biol. Macromol.*, 2019, **133**, 382–390.
- 52 W. Xu, J. Xia, S. Min and Y. Xiong, *Spectrochim. Acta, Part A*, 2022, **274**, 121034.
- 53 B. A. McGregor, X. Liu and X. G. Wang, *J. Text. Inst.*, 2018, **109**, 813–822.
- 54 L. C. Lee, C.-Y. Liong and A. A. Jemain, *Analyst*, 2018, **143**, 3526–3539.
- 55 R. G. Brereton, *Chemometrics for Pattern Recognition*, John Wiley & Sons Ltd., Chichester, England, 2009.
- 56 L. C. Soares, J. O. Alves, L. A. Linhares, F. B. E. Filho and M. P. F. Fontes, *Microchem. J.*, 2017, **133**, 258–264.



- 57 J. Trevisan, P. P. Angelov, P. L. Carmichael, A. D. Scott and F. L. Martin, *Analyst*, 2012, **137**, 2302–2312.
- 58 L. Wu, B. Du, Y. V. Heyden, L. Chen, L. Zhao, M. Wang and X. Xue, *Trac, Trends Anal. Chem.*, 2017, **86**, 25–38.
- 59 Y. Shai, *Biochim. Biophys. Acta*, 2013, **1828**, 2306–2313.
- 60 S. Saravanan, D. K. Sameera, A. Moorthi and N. Selvamurugan, *Int. J. Biol. Macromol.*, 2013, **62**, 481–486.
- 61 B. C. Powell and G. E. Rogers, *Hair Keratin: Composition, Structure and Biogenesis*, Springer, Berlin, Heidelberg, 1986, pp. 695–721.
- 62 Z. Ganim, H. S. Chung, A. W Smith, L. P Deflores, K. C. Jones and A. Tokmakoff, *Acc. Chem. Res.*, 2008, **41**, 432–441.
- 63 Á. I. López-Lorente and B. Mizaikoff, *Anal. Bioanal. Chem.*, 2016, **408**, 2875–2889.

