

Detection of goat body fat adulteration in pure ghee using ATR-FTIR spectroscopy coupled with chemometric strategy

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Abstract Ghee forms an important component of the diet of human beings due to its rich flavor and high nutritive value. This high priced fat is prone to adulteration with cheaper fats. ATR-FTIR spectroscopy coupled with chemometrics was applied for determining the presence of goat body fat in ghee (@1, 3, 5, 10, 15 and 20% level in the laboratory made/spiked samples). The spectra of pure (ghee and goat body fat) and spiked samples were taken in the wavenumber range of 4000–500 cm⁻¹. Separated clusters of pure ghee and spiked samples were obtained on applying principal component analysis at 5% level of significance in the selected wavenumber range (1786–1680, 1490–919 and 1260–1040 cm⁻¹). SIMCA was applied for classification of samples and pure ghee showed 100% classification efficiency. The value of R² was found to be >0.99 for calibration and validation sets using partial least square method at all the selected wavenumber range which

indicate that the model was well developed. The study revealed that the spiked samples of goat body fat could be detected even at 1% level in ghee.

Keywords Pure mixed ghee · Goat body fat · Spike · ATR-FTIR · Principal component analysis · Partial least square

Introduction

Milk fat is one of the most important dietary components for supplying nutrients. The composition of milk fat is unique as it contains various bioactive components like butyric acid, cis and trans palmitoleic acid, phytanic acid, conjugated linoleic acid (CLA), alpha-linolenic acid (ALA) which are potentially identified as positive predisposing factors for human health (Belury 2002; McBain et al. 1997; Parodi 1994). Butyric acid and linolenic acid are reported to have anticarcinogenic properties. Linolenic acid is also shown to prevent hypertension and heart related diseases and is involved in the improvement of vision (Parodi 2003). CLA is reported to induce antithrombotic effect (Truitt et al. 1999), enhance immunological functions (Pariza et al. 2001), decrease total and low density lipoprotein (LDL) cholesterol (Lee et al. 1994) and inhibit carcinogenesis (Ip et al. 1999). Ghee (heat clarified milk fat) is reported to be the richest natural source of CLA (Sserunjogi et al. 1998). Most dietary guidelines recommend the consumption of milk and dairy products as an important part of a healthy well-balanced diet (Gidding et al. 2009 and United States Department of Agriculture Department of Health and Human Services 2010). Milk fat is valued higher over other fats and oils. Report from National Sample Survey Office: Level and Pattern of Consumer Expenditure (2011–2012) revealed that higher

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income brackets spend more on dairy products (Mani and Intodia 2014).

India is the world's largest milk producing nation. However, there is a gap between the demand and supply as the growth in milk production is 3.3%, while there is 5% growth in the consumption (<http://www.dairyuniverseindia.com/Marketdate.html>) (last accessed on 19th June, 2015). Similarly, the supply of ghee (most consumed after fluid milk among dairy products), also falls short of its demand (Das 2012; Mani and Intodia 2014). The higher cost of ghee (4–5 times) over the other edible fats and oils and the limited supply of this high premium fat attract the unscrupulous manufacturers to adulterate it with cheaper alternatives, especially in the Indian subcontinent. A wide range of literature is available on the incidences of adulteration of ghee with foreign fats (Gandhi 2009; Headlines Today 2009; Ramkumar 2014). Although, this malpractice of adulterating the ghee is not new. A statistical survey of 13 years, 1960–1972, on ghee revealed that 16% of the samples investigated were adulterated, with average level of adulteration being 12%. The more recent reports suggest a higher percent of adulteration in the ghee samples. Gahlawat et al. (2012) reported that all of the ghee samples collected from Delhi region, India contained vanaspati (hydrogenated vegetable oils) as an adulterant. In another survey, Amrutha Kala (2013) reported that up to 43% branded and 87.5% of unbranded ghee samples did not conform to triglyceride profiles of pure ghee based on triglyceride profile and standardized values. Hence, such adulteration is of major concern as it leads to fraud by overcharging (Upadhyay et al. 2015). This practice can affect the nutritional value of original product, its technological properties and most importantly it violates the consumer rights (Nunes 2014). By adding animal body fats in ghee, one of the sacred foods, the oblivious ghee traders lacking honesty not only fleece the public, but also play with their religious sentiments, especially of the devourer and vegetarian sect of the society. The consumption of animal body fat (from red meat) is reported to have adverse health implications. Willett et al. (1990) and Giovannucci and Willett (1994) reported that its consumption increase the risk of colon cancer. Trans fatty acids are reported to be present in animal body fat and are associated with deleterious effects on serum lipoproteins leading to increase in the risk of cardiovascular and cerebrovascular diseases (Stender and Dyerberg 2004).

The detection of adulteration of ghee with animal body fat is one of the priority areas of research in the field of food technology as only limited methods are available for their detection. The available tests are mainly based on planar chromatography technique (Ramachandra and Dastur 1959; Chakrabarty et al. 1968) that involves hazardous chemicals; determination of fatty acid composition of

triglycerides and 2-monoglycerides (Soliman and Younes 1986). But, these tests suffer from repeatability problems or do not give clear cut range of determination of adulteration. The other methods are based on techniques like differential scanning calorimetry (Lambelet and Ganguli 1983; Coni et al. 1994) which are reported to detect the presence of foreign animal body fat in ghee only at the higher levels.

In the light of above facts, there is an increased interest of researchers in finding a solution to the problem of detection of adulteration of animal body fat in ghee. Fourier transform infrared (FTIR) spectroscopy, a fingerprint technique, is rapid, simple, sensitive, advanced and powerful analytical tool that finds its application in the wide areas of detection of adulteration in various food commodities (Sivakesava and Irudayaraj 2001; Ozen and Mauer 2002; Jha and Matsuoka 2004; Rodriguez-Saona and Allendorf 2011). FTIR spectroscopy is also reported to provide information about the composition of dairy product from the complex spectra (Albanell et al. 1998; Sørensen and Jepsen 1998; Rodriguez-Otero et al. 1997; Lerma-García et al. 2010; Nicolaou et al. 2010). The technique has been widely applied, because once the instrument has been calibrated; it can be used for routine analyses. FTIR coupled with chemometric techniques had been extensively used to discriminate different edible oils and fats (Safar et al. 1994; Rohman and Man 2010; Nurrullhidayah et al. 2013). However, no work has been done to distinguish ghee from goat body fat using FTIR spectroscopy. Thus, present study was aimed with the objective of detection of adulteration of goat body fat in pure mixed ghee (heat clarified milk fat) by employing FTIR spectroscopy coupled with chemometric techniques.

Materials and methods

Collection of Sample

Pure cow milk and pure buffalo milk were separately collected from their respective pooled milk from the Institute Farm, Guru Angad Dev Veterinary and Animal Sciences University, Ludhiana, Punjab (India). The goat adipose tissues were separately procured from the local slaughterhouse of Ludhiana, Punjab (India).

Preparation of sample

Pure ghee from cow milk and buffalo milk were prepared by employing direct cream method (De 2012). The ghee thus obtained was subjected to filtration using four folds of the filter paper to get clear ghee devoid of ghee residues. The two ghee vis-à-vis pure cow ghee and pure buffalo

ghee were mixed together in the ratio of 1:1 to get the pure mixed ghee (PMG). The samples were stored in glass bottles (of Borosil make) under refrigerated conditions (4–6 °C) until further use. Goat body fat (GBF) was prepared from the adipose tissue procured from the local slaughter house by dry rendering process at 152 ± 2 °C till the complete extraction of fat (Upadhyay et al. 2014). For preparation of adulterated samples, GBF was spiked into PMG at a level of 1, 3, 5, 10, 15 and 20% to make the binary system comprising of PMG and GBF. Infrared spectra were collected for each sample.

Spectra Acquisition

The fats and oils are mainly composed of triglycerides and can be directly applied in their neat form to ATR crystal (Nurrulhidayah et al. 2013). Thus, using an auto-pipette, fat samples obtained were placed in direct contact with Diamond crystal cell attenuated total reflectance (ATR) crystal (Path length: 1.66 μm ; 7.2 Build, Bruker, Model-Alpha, Germany). The absorption spectra of the samples were acquired using Fourier transform infrared (FTIR) spectrometer at resolution of 4 cm^{-1} and scan speed of 0.2 cm s^{-1} . Twenty four scans of each sample maintained at 70 °C (so as to maintain the sample in completely molten stage) were taken in the wavenumber range of $4000\text{--}500\text{ cm}^{-1}$ and connected to software OPUS (7.2 Build, Bruker Optik GmbH) for collection of data. The reference (background) spectra using blank ATR crystal were recorded at an interval of 35 spectra of 24 scans each. The reference spectra were used by the instrument in rationing the spectra of the samples in the absorbance unit. After each scan, the ATR crystal was carefully cleaned by wiping in situ with a soft tissue paper using isopropanol and dried completely before taking the next spectra of the sample.

Chemometric analysis

All the spectra were exported to the Unscrambler software (version 10.2; CAMO AS, Trondheim, Norway) and divided into two groups of calibration (training) and validation (testing) sets, taking every third sample as validation set. This amounted to be around 187 and 93 samples in respectively, calibration and validation sets of the PMG adulterated with GBF. Care was taken that control (PMG) and spiked samples (PMG containing GBF) were well represented in calibration and validation sets to avoid the confusion of calibration samples in validation sets. Absorbing regions of the spectra dictate the number of variables used as input in the latent variables based methods such as Principal Component Analysis (PCA) and partial least square (PLS) regression modeling. As a thumb

rule, the spectral range should include regions where information pertaining to variation in the concentration of analyte to be modeled and any other matrix constituents are well represented. However, the spectral range should exclude regions that are dominated by artifacts or noise. The raw spectra were plotted and peaks were looked into them so as to identify and select the suitable region. PCA was performed so as to examine the clustering of samples in different groups and also for the purpose of identifying the outliers, if any.

Classification of samples

Initially a global model was built followed by development of class model for control and spiked samples using calibration set of samples. Soft Independent Modeling of Class Analogy (SIMCA) thereafter was used to predict the class memberships using validation set of samples at 5% level of significance. Test samples were classified to one of the established class models on the basis of their best fit to the respective model. False negative and false positive samples were identified to calculate the classification efficiencies. The samples which belonged to a specific class but did not classify as such by the class model were referred to as false negative, while those samples which were incorrectly identified in the relevant class model but did not belong to that specific class were referred to as false positive.

Validation of results

PLS regression using full cross validation option present in the Unscrambler software was performed on the peaks and depressions of the selected spectral window (1786–1680, 1490–919 and $1260\text{--}1040\text{ cm}^{-1}$) for developing the model for predicting spiked GBF in PMG. The best model was selected based on the standard error of calibration (SEC), coefficient of determination (R^2) and standard error of prediction (SEP) (Jaiswal et al. 2012; Jha and Matsuoka 2004; Jha et al. 2015).

Results and discussion

Natural fats contain a wide range of triacylglycerol species with fatty acids of different substitution patterns, chain length, degree of unsaturation and other minor components. Milk fat is one of the unique naturally occurring fats of animal origin containing almost 70% of saturation of which around 11% comprises of short-chain fatty acids, approximately half of which is butyric acid. Around 25% of the fatty acids in milk fat are mono-unsaturated, while 2.3% are poly-unsaturated (Månsson 2008). Goat depot fats are reported to contain around 55% saturated fatty acid,

39% monounsaturated fatty acid and 4% polyunsaturated fatty acid (Casey and Van Niekerk 1985). Although the type of fatty acids present in different fats was more or less same but the concentration of these varies. In past, few researchers attempted to use Ultraviolet spectra for differentiation of pure ghee from the ghee containing animal body fat. Singhal (1973) and Sharma (1989) dissolved cow ghee, buffalo ghee and animal body fats (buffalo, goat, pig and sheep) in n-hexane and scanned the Ultraviolet spectra of these between 200 and 320 nm. They reported a maximum absorption between 220 to 230 nm. However, Kumar et al. (2010) and Sharma (1989) reported that adulterated ghee could not be differentiated from pure ghee using the above method. On the contrary, FTIR can be considered a fingerprint technique as no two fats and oils have superimposing FTIR spectra and specific functional groups show specific absorption maxima (Bendini et al. 2007).

FTIR spectral analysis

The representative FTIR spectra of control, spiked and GBF samples in the region of 4000–500 cm^{-1} is shown in Fig 1 in Electronic Supplementary Material (ESM). It revealed that the peak height for PMG and GBF were non-overlapping, while the peaks of the spiked samples fell between the two as per the level of spiking in entire range of the region of the spectra. The peaks of samples that had more percentage of spiking were observed to be near to peak of GBF, while the samples that contained lower percentage of spiking were near to PMG. However, the differences were more distinct in the wavenumber regions of 1786–1680 and 1490–919 cm^{-1} (Fig. 1). The peaks in the smaller spectral range of 1260–1040 cm^{-1} also showed distinct differences among the pure and spiked samples of GBF in PMG. This could be due to the differences in the biochemical composition of PMG and GBF as explained below.

The absorptions in the FTIR spectroscopy were attributed to the particular spectrum absorbed by the specific functional group. For example, the functional group associated with $\text{C}=\text{O}$ (ester) i.e. carbonyl group from the ester linkage of triacylglycerol is attributed to the wavenumber of 1745 cm^{-1} . The wavenumbers 1460, 1377, 1161, 1061, 1117 and 1097 cm^{-1} have been reported to be associated with different functional groups/bonds found in the fatty acids i.e. $\text{C}-\text{H}-(\text{CH}_2, \text{CH}_3)$, $=\text{C}-\text{H}-(\text{cis})$, $\text{C}-\text{O}-\text{CH}_2-$, $\text{C}-\text{H}-$ and $\text{C}-\text{H}-$ bending, respectively (Nurrulhidayah et al. 2010; Vlachos et al. 2006; Guillén and Cabo 1997; Safar et al. 1994). Therefore, in the present study, the peaks observed in the region of 1786–1680 cm^{-1} might be attributed to the ester group of the triglycerides present in the fat samples, while the peaks observed in the spectral region of 1490–919 cm^{-1} might be associated with the

different functional groups present in the fatty acid. Earlier workers, Guillén and Cabo (1997) suggested bands near 1400 and 1097 cm^{-1} as the fingerprint regions, while Nurrulhidayah et al. (2013) reported a wider range of 1500–1000 cm^{-1} as the fingerprint regions for detection of beef fat in butter.

Principal component analysis

Principal Component Analysis (PCA) used the combinations of absorbance at the specified wavenumbers that explained most of the variance present in the data set. PCA of the spectral data carried out in the selected range of wavenumber i.e. 1786–1680 cm^{-1} so as to observe the clustering of samples for the control, spiked and GBF samples is represented in Fig. 2a. It can be observed from the figure that PMG and GBF showed well separated clusters. The variance as explained by Principal Component 1 and 2 were 91% and 9%, respectively indicating the role of FTIR in the detection of GBF in PMG at 5% level of significance. With the increase in the level of spiking, the clusters of the spiked samples moved away from the cluster of PMG and approached to the cluster of GBF. In fact, even the lowest level of spiking showed separate clusters of the sample than the PMG. This could again be explained by the difference in the fatty acid composition of pure mixed ghee and goat body fat. Similar results were obtained when the PCA was applied to the other range of the selected wavenumbers i.e. 1490–919 cm^{-1} and 1260–1040 cm^{-1} (Fig. 2b, c), where the Principal Component 1 and 2 contributed respectively, 92 and 8% and 97 and 3%. Recently, Nicolaou et al. (2010) used FTIR spectroscopy and multivariate analysis for detection of presence of sheep and goat milk in cow milk. However, in the present investigation, FTIR coupled with chemometrics was used for detection of goat body fat in pure mixed ghee with the purpose of detection of adulteration of cheaper body fat in the high priced mixed ghee.

Classification of sample

Soft Independent Modeling of Class Analogy (SIMCA) approach at the selected wavelengths was adopted for predicting probable class membership among the PMG, PMG spiked with GBF at different levels and GBF. As a result of SIMCA class projections of the spectra, it was evident that PMG was always classified under its class with the classification efficiency of 100% at all the selected wavenumbers (Table 1). Similarly, GBF was always classified in its own class at all the wavenumber ranges studied giving a classification efficiency of 100%. It was further interesting to note that although the spiked samples were

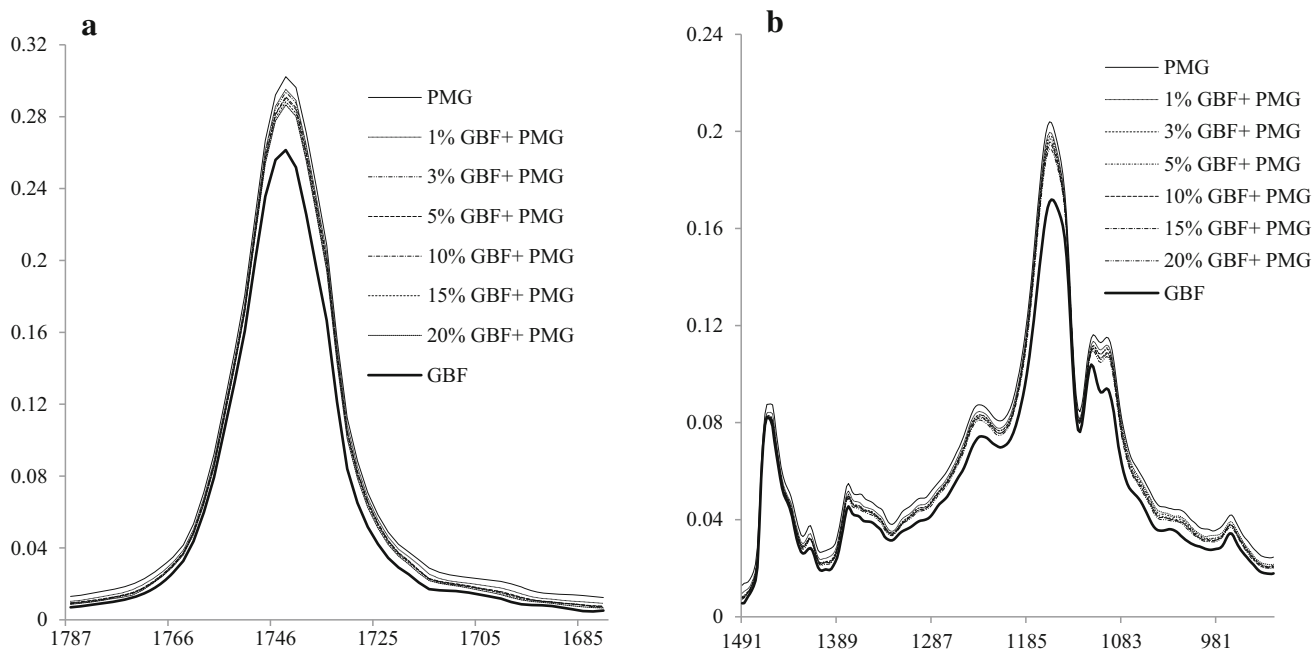


Fig. 1 Spectra of mixed ghee along with the addition of goat body fat in the wavenumber range **a** 1786–1680 cm^{-1} , **b** 1490–919 cm^{-1} . *PMG* pure mixed ghee (1:1—pure cow ghee and pure buffalo ghee), *GBF* goat body fat

misclassified at times but they never classified in the PMG indicating the scope of the methodology for detecting the presence of GBF in PMG. Minimum classification efficiency for the test samples (those spiked with GBF) was obtained to be 90.90% for 1490–919 and 1260–1040 cm^{-1} spectral ranges for the samples spiked with up to 5% of the GBF. However, in case of 1786–1680 cm^{-1} spectral range, the minimum class efficiency was observed to be 92.67% for the samples spiked with up to 3% of GBF in PMG. However, the classification efficiency was 100% at the higher levels. Thus, it can be concluded that even the lowest level of 1% GBF in PMG can be detected using SIMCA approach considering the results that none of the spiked samples were classified in the PMG class.

Calibration and validation

Partial least square (PLS) was used for constructing calibration model in selected ranges of wavenumbers and the effect of different spectral windows on spectral data modeling are presented (Table 2). The results revealed R^2 closer to unity (i.e. >0.99) in all the selected spectral ranges, which further signify the better relationship between actual and FTIR predicted values of the analyte of interest (GBF in our case). Also, according to International Conference on Harmonization (ICH 1994), R^2 value higher than 0.99 is acceptable for such relationship which was observed in all the selected wavelengths. RMSEC refers to the calibration uncertainty. The difference between

RMSEC and RMSEV was less, which indicate that the calibration model was accurately developed. Also, the slope was observed to be around 45° , while the intercept was negligible for the scatter plots of both calibration and validation sets of sample (Fig. 2 in ESM) confirming the predictability of the analyte i.e. GBF in PMG. It can also be said that all the selected wavenumber ranges (1786–1680, 1490–919 and 1260–1040 cm^{-1}) are useful for determining the presence of GBF in PMG as these are associated with the specific groups present in the fat as mentioned previously. Infra-red spectroscopy was also used by some workers (Konevets et al. 1987; Sharma 1989) in the past who reported a level of detection of 10% for the animal body fats in milk fat, which is much higher than the one reported in the present study.

Conclusion

The difference in the absorption values of pure mixed ghee, goat body fat and the pure mixed ghee spiked with goat body fat in the spectral region of 1786–1680, 1490–919 and 1260–1040 cm^{-1} is clearly evident from the peaks in the said regions. Using SIMCA, almost 97% of the test samples could accurately classify themselves into their respective class in the models developed without preprocessing the data. The efficiency of the coefficient of determination using PLS for quantitatively predicting the data was also found to be high for both calibration and

Fig. 2 Principal component scores plot depicting clusters of pure mixed ghee, pure goat body fat and mixed ghee with different levels of goat body fat in the wavenumber range of **a** 1786–1680 cm^{-1} , **b** 1490–919 cm^{-1} , **c** 1260–1040 cm^{-1}

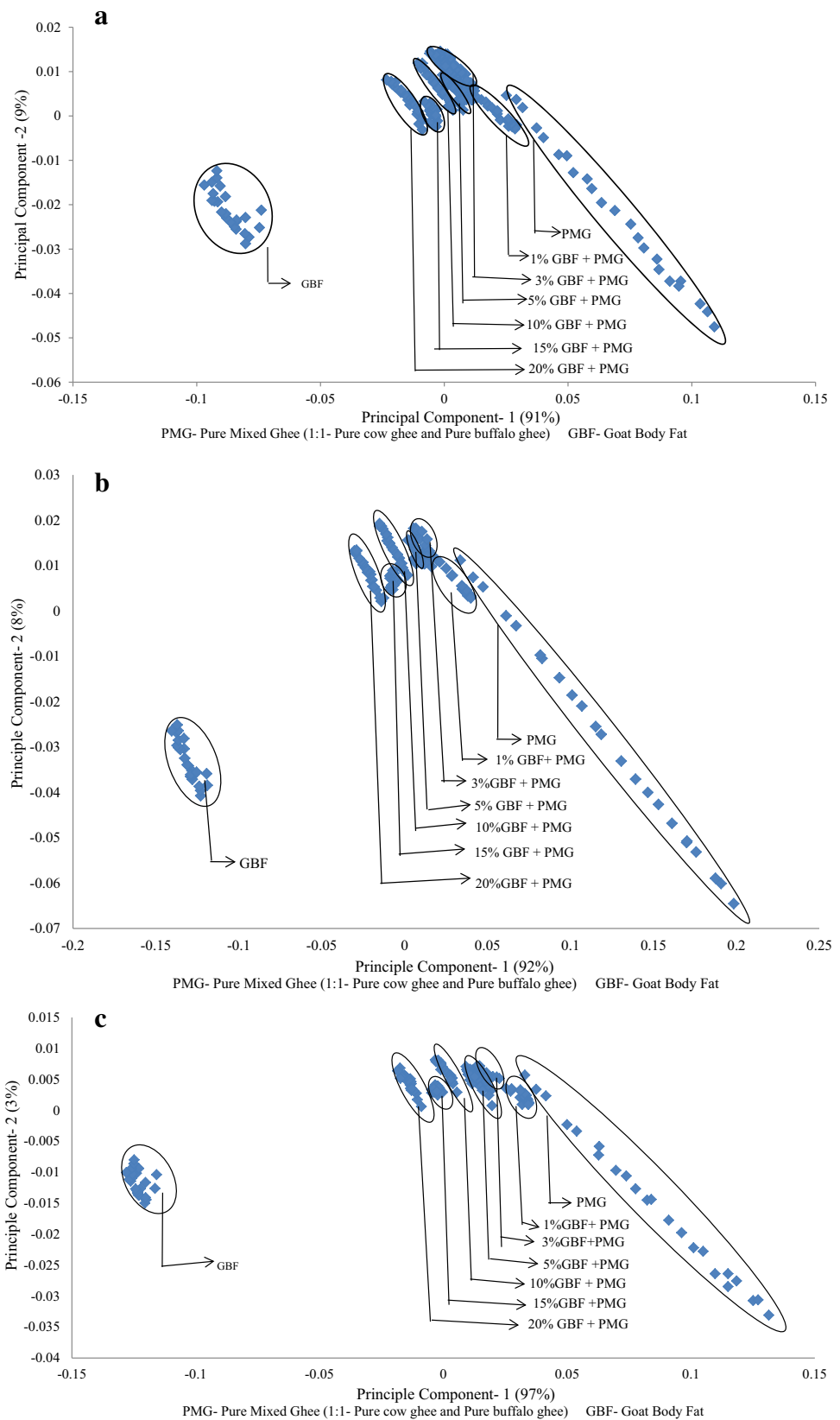


Table 1 SIMCA classification results for mixed ghee along with the addition of goat body fat

Wavenumber (cm ⁻¹)	% GBF in PMG	Total no. of samples	Number of selected classes								Misclassified	Classification efficiency (%)
			0	1	3	5	10	15	20	100		
1786–1680	0	12	12	–	–	–	–	–	–	–	0	100
	1	12	–	11	–	–	–	–	–	–	1	92.67
	3	12	–	–	11	1	–	–	–	–	2	92.67
	5	11	–	–	3	11	–	–	–	–	3	100
	10	12	–	–	–	–	12	–	–	–	0	100
	15	12	–	–	–	–	–	12	–	–	0	100
	20	11	–	–	–	–	–	–	11	–	0	100
	100	12	–	–	–	–	–	–	–	12	0	100
1490–919	0	12	12	–	–	–	–	–	–	–	0	100
	1	12	–	11	–	–	–	–	–	–	1	92.67
	3	12	–	–	11	–	–	–	–	–	1	92.67
	5	11	–	–	–	10	–	–	–	–	1	90.90
	10	12	–	–	–	–	12	–	–	–	0	100
	15	12	–	–	–	–	–	12	–	–	0	100
	20	11	–	–	–	–	–	–	11	–	0	100
	100	12	–	–	–	–	–	–	–	12	0	100
1260–1040	0	12	12	–	–	–	–	–	–	–	0	100
	1	12	8	11	–	–	–	–	–	–	9	92.67
	3	12	–	–	11	1	–	–	–	–	2	92.67
	5	11	–	–	2	10	–	–	–	–	3	90.90
	10	12	–	–	–	–	12	–	–	–	0	100
	15	12	–	–	–	–	–	12	–	–	0	100
	20	11	–	–	–	–	–	–	11	–	0	100
	100	12	–	–	–	–	–	–	–	12	0	100

Table 2 Effect of different spectral windows on spectral data modeling using PLS for the selected range of wavenumbers for detection of adulteration of goat body fat in pure ghee

Wave number range (cm ⁻¹)	Calibration			Validation		
	R ²	RMSEC	Bias	R ²	RMSEV	Bias
1786–1680	0.993	2.586	0.0	0.993	2.625	–0.009
1490–919	0.997	1.583	0.0	0.997	1.615	–0.005
1260–1040	0.997	1.636	0.0	0.997	1.687	–0.003

validation sets (>0.99) at all the wavelengths studied (1786–1680, 1490–919 and 1260–1040 cm⁻¹). The difference in the value of RMSEC and RMSEV was less, indicating that the model was well developed. The study concludes that the method is rapid, non-destructive and has great precision and accuracy. Attenuated Total Reflectance- Fourier Transform Infra-red spectroscopy has the potential of determining the adulteration of even 1% of goat body fat in pure mixed ghee.

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Compliance with ethical standards

Conflict of interest The authors do not have any conflict of interest.

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