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Development of new NIR-spectroscopy method combined with multivariate analysis for detection of adulteration in camel milk with goat milk

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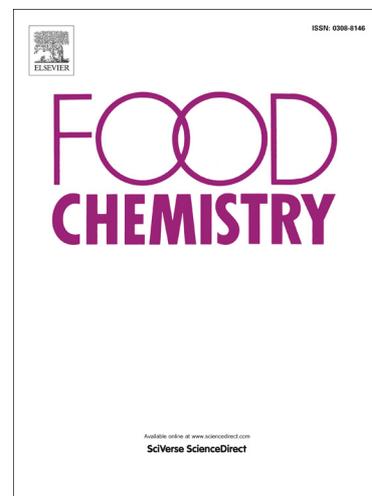
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31 model was used to quantify the levels of adulteration from, 0%, 2%, 5%, 10%, 15%
32 and 20%. The PLS model showed the RMSEC = 1.10% with $R^2 = 94\%$. This method
33 is simple, reproducible, having excellent sensitivity. The limit of detection was found
34 0.5 %, while the limit of quantification was 2 %.

35 **Keywords:**

36 NIR- spectroscopy; Camel milk adulteration; PCA, PLS-DA, PLS regression

37

38 **Introduction:**

39 In many arid and semi-arid areas of the world camel is of considerable socio-
40 economic value and its milk has a number of nutrients beneficial to human body
41 (Shamsia, 2009). For centuries, people have been using camel for transportation and
42 camel milk as a source of food and also as medicines for several diseases (Gizachew,
43 Teha, & Birhanu, 2014). In addition, it has been consumed for centuries due to its
44 nutritional values and medicinal properties (Dowelmadina, Zubeir, Arabi, & Abaker,
45 2015). In the last twenty years, there have been many studies on the use of camel milk
46 in the treatment of human diseases (Kaskous, 2015).

47 For both water-soluble and fat-soluble vitamins, milk is a valuable source. Because of
48 its high concentration of vitamin C, camel milk is a kind of exception. Camel milk has
49 30 times more Vitamin C than cow milks, and contains 6 times more than human
50 milk. In the desert areas this is highly important, where vegetables and fruits are
51 scarce. Therefore, in the diet of inhabitants of these regions camel milk is often the
52 only source of vitamin C (Gizachew, Teha, & Birhanu, 2014). The composition of
53 camel milk is different from other ruminants' milk. It has low sugar, low cholesterol,
54 high minerals (potassium, sodium, copper, iron, magnesium and zinc), low protein,
55 high vitamins, and high concentrations of insulin (Yadav, Kumar, Priyadarshini, &
56 Singh, 2015).

57 Being a rich source of the nutritional constituents and variety of biological activities
58 that effect development and growth of specific body organs, metabolic responses
59 towards nutrients absorption, digestion and fight against diseases. By the digestive
60 action on milk biologically active peptides are produced in the gastrointestinal tract.

61 The positive health effects of milk proteins can be presented as anti-microbial,
62 antioxidative, anti-thrombotic, and antihypertensive or immunomodulatory
63 (Antanasova & Ivanova, 2010).

64 Camel's milk is considered as abundant source of protein for people living in arid
65 lands of the world. This protein is rich in protective components including lysozyme,
66 lactoferrin, Lactoperoxidase (LPS), and peptidoglycan recognition protein (PGRP)
67 which are found only in camel's milk (Singh, Ghorui, & Sahani, 2006), IGA and IGg
68 immunoglobulins that are suitable with human ones and provide effective defense
69 against several viral and bacterial pathogens. The fact that camel's milk is low in
70 different β -caseins (Beg, von Bahr-Lindstrom, Zaidi, & Jornvall, 1986) and without
71 β -lactoglobulin (Merin et al., 2001) makes it more attractive for those suffering from
72 milk allergies (Makinen-Kiljunen & Palosuo, 1992; Shabo, Brazel, Margoulis, &
73 Yagil, 2005).

74 Milk is an important source of mineral substances, especially sodium, potassium,
75 calcium, magnesium, phosphorus, chloride, iodine, and small amounts of iron. The
76 key mineral compounds of milk are phosphorus and calcium, which are significant for
77 the proper development of new borns and bone growth. The high bioavailability of
78 these minerals effects the distinctive nutritional value of milk. Camel milk is the
79 richest in these minerals. The mean values in mineral contents of dromedary camel
80 milk (100g-1) for zinc (0.53 mg), manganese (0.05 mg), magnesium (10.5 mg), iron
81 (0.29 mg), sodium (59 mg), potassium (156 mg) and calcium (114 mg) (Elamin, &
82 Wilcox, 1992; Sawaya, Khalil, Al-Shalhat, & Al-Mohammad, 1984).

83 According to the previous studies, camel milk possesses nutritional and medicinal
84 values (Gizachew, Teha, & Birhanu, 2014). It is rarely available in the market and
85 sold with much higher prices, and thus called as desert gold because of its higher
86 prices. Its adulteration with goat, cow, buffalo and other inexpensive commercial
87 milks has already been started. In the present study, the focus is to develop a new NIR
88 spectroscopy method combined with chemo metrics to authenticate as well as to
89 check the level of adulteration in camel milk with goat milk as an adulterant. The
90 novelty of the method lies in high sensitivity, better reproducibility, more economical,
91 and less or no sample preparations for the overall experiment.

92 **2. Materials and methods**

93 **2.1. Camel milk Samples preparation**

94 Three different camel milk as well as goat milk samples were collected from
 95 Aldhahira as well as Sharqia regions of Sultanate of Oman and analysed. Those camel
 96 milk samples were then adulterated with goat milk at five different percentage levels:
 97 2, 5, 10, 15 and 20 %. Number of total samples used was 54: 9 pure (3x3=9), 45
 98 adulterated with goat milk. The samples were prepared in triplicate. All the samples
 99 were joined together and split into two sets (70 % of the samples) training set and (30
 100 % of the samples) a test set for validation for PLS regression,.

101 **2.2. Apparatus**

102 Perkin Elmer Frontier NIR spectrophotometer was used to measure all the samples in
 103 absorption mode in the wavelength range from (700-2500 nm), at 2 cm⁻¹ resolution
 104 and using a 0.2 mm path length CaF₂ sealed cell. Prominent absorption peaks were
 105 appeared in the region from 4000 to 7500 cm⁻¹ wavenumber.

106
 107 **2.3. Statistical analysis**

108 For statistical analysis the Unscrambler version 9.0 by Camo and Microsoft Excel
 109 2010 were used The PLS-DA, PCA, and PLS regression models were built for both
 110 pure and adulterated camel milk samples. Spectral pre-treatments, such as 1st
 111 derivative with Savitzky-Golay with 5 smoothing point 2nd order polynomial and
 112 baseline correction were conceded. To validate the PLS-DA models Leave-one-out
 113 cross validation was used. For PLS regression all the samples were joined together
 114 and split into two sets, (70% of the samples) a training set and (30% of the samples) a
 115 test set for validation. To validate the PLS regression models built with the training
 116 set, external cross validation was used. The RMSECV (Root Mean Square Error of
 117 Cross Validation) was used as an internal indicator of the predictive ability of the
 118 models. Using Eq. 1: RMSECV is calculated as:

119

$$120 \quad RMSECV = \sqrt{\frac{\sum_{i=1}^n (y_i - \hat{y}_i)^2}{n}} \text{-----(1)}$$

121

122 Where \hat{y}_i is the % of adulteration predicted by the model, y_i is the actual % of
 123 adulteration (measured value) and n is the number of segments left-out in the cross-
 124 validation method, which is equivalent to the number of samples of the training set.
 125 RMSECV smaller values are the indicative of a improved prediction ability of the
 126 model.

127

128 The statistical measure RMSEP is used to tell how well the model predicts new
 129 samples (not used when building the model) which is calculated using Eq. 2:

130

$$RMSEP = \sqrt{\frac{\sum_{i=1}^{n_t} (y_{t,i} - \hat{y}_{t,i})^2}{n_t}} \quad \text{---(2)}$$

131

132 where $\hat{y}_{t,i}$ is the % of adulteration predicted by the model, $y_{t,i}$ is the measured value
 133 (actual % of adulteration), and n_t is the number of samples in the test set. Average
 134 error expresses by RMSEP to be expected in future predictions when to the unknown
 135 samples the calibration model is applied.

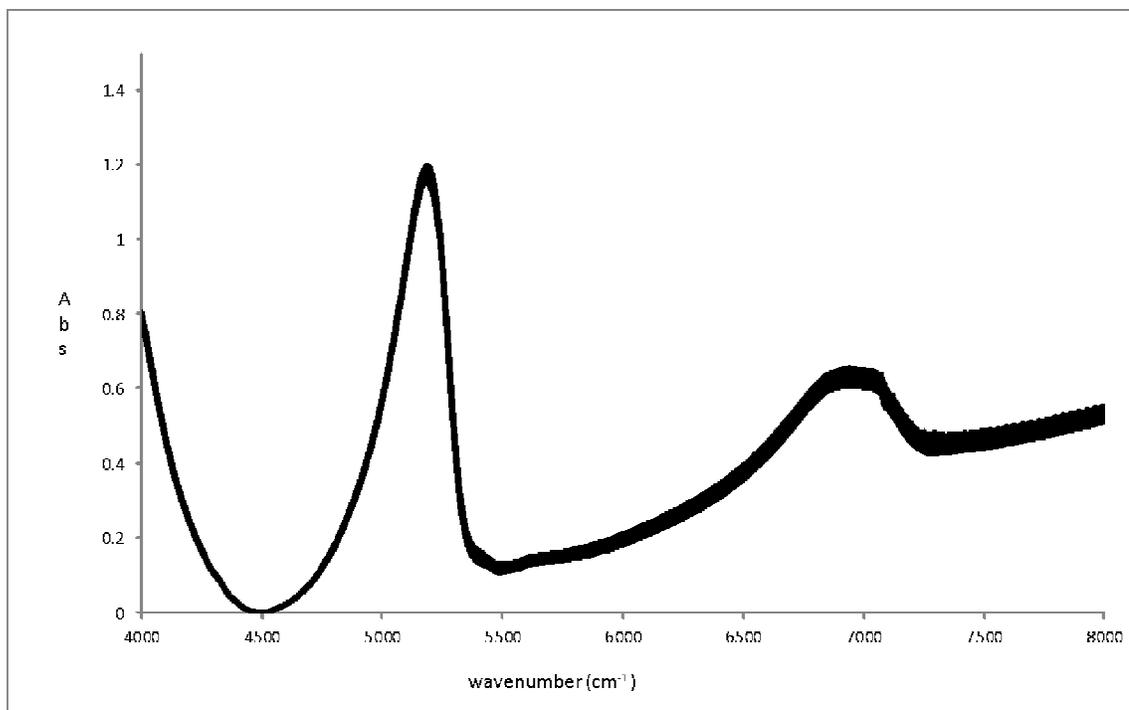
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137 3. Results and discussion:

138 3.1. Near Infrared spectra

139 Figure 1 shows the spectra of NIR for all samples ranging from (10000-4000 cm^{-1}) in
 140 term of wavenumbers while in term of wavelength ranging from (700-2500 nm) using
 141 a 0.2mm path length CaF_2 sealed cell.

142



143

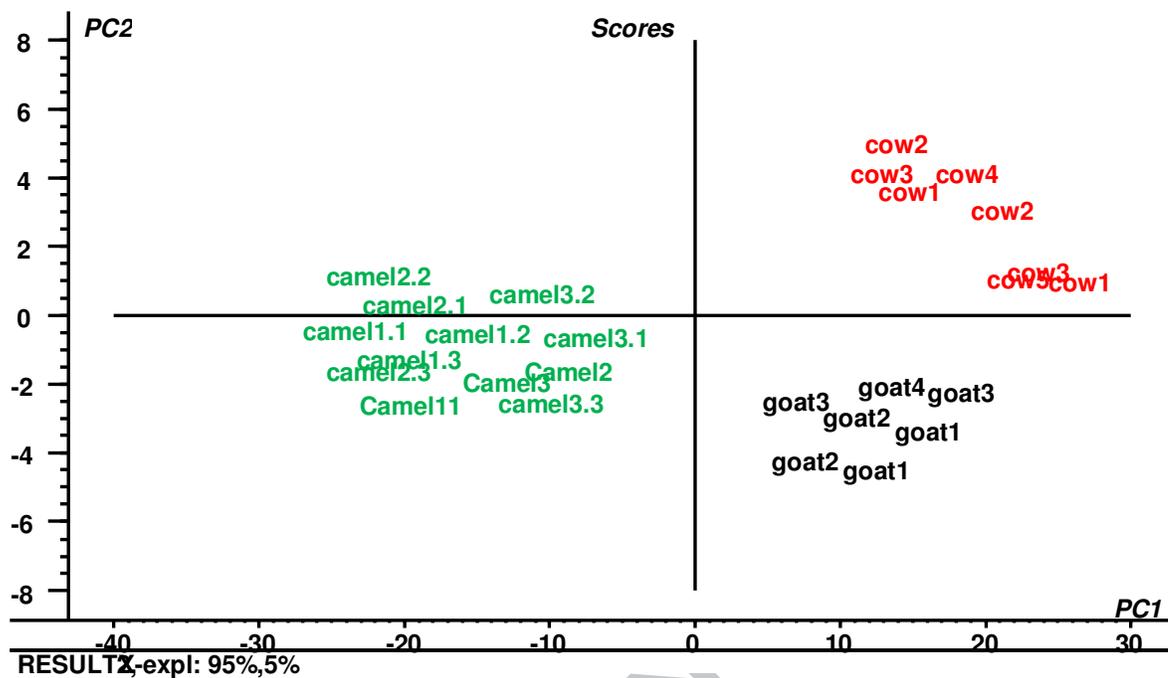
144 Figure 1. NIR spectra for milk samples, (point spectra) pure camel milk and (solid
145 line spectra) adulterated with goat milk.

146

147 Figure 1 shows the NIR spectra for both pure camel milk (point spectra) and
148 adulterated with goat milk (solid line spectra). It can be seen from the that there is a
149 prominent absorption peaks at wavenumber 5198 cm^{-1} .

150 Although the spectra appear to be very similar. In order to see the difference among
151 the unadulterated camel, cow and goat milk an alternative approach of principal
152 componentes analysis (PCA), was applied in which a PCA model was built as shown
153 in Figure 2. It can be seen from the PCA plot that there is complete differentiation and
154 separation among the camel, cow and goat milk. They are spaced and grouped in the
155 specific different regions of the PCA plot.

156



157

158 Figure 2. PCA plot for the unadulterated camel, cow and goat milk

159

160 Similarly in order to detect the discrimination between the pure and adulterated
 161 camel milk samples, PLS-DA (Partial least-squares discriminant analysis) method was
 162 applied. It was built for the spectral data between pure and with 10 % goat milk
 163 adulteration level as shown in Figure 3. Spectral pre-treatments, such as baseline
 164 correction, and Savitzky-Golay smoothing were used for building this model.

165

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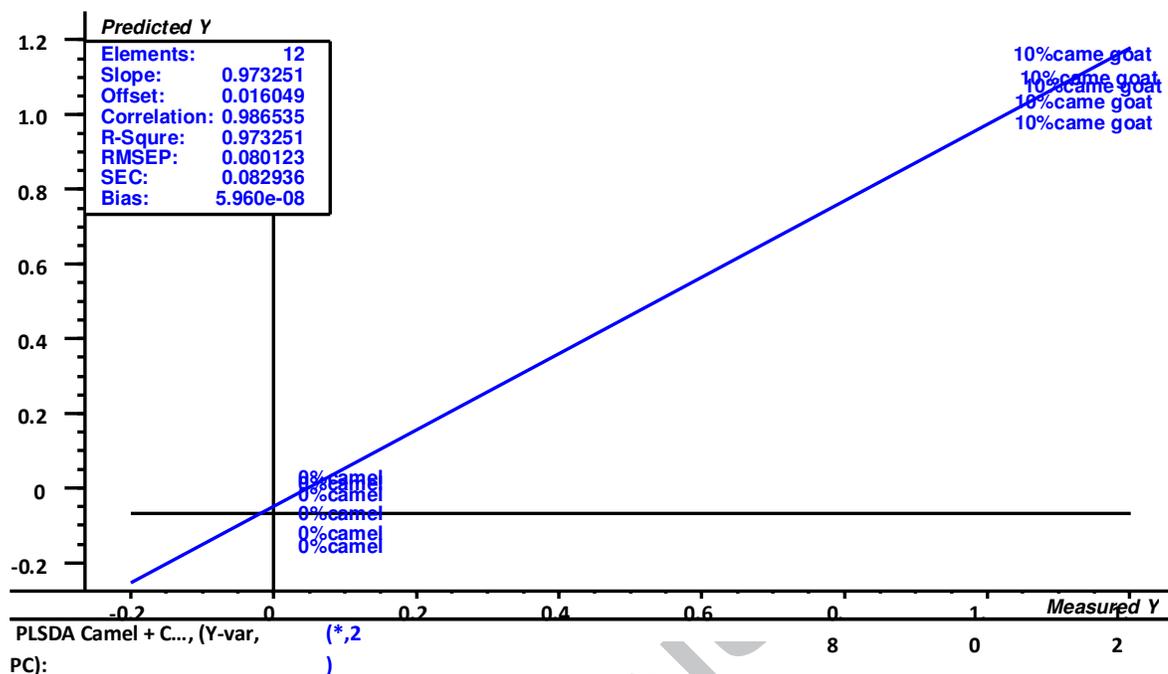
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173 Figure 3. PLS-DA model for pure camel milk and with 5% goat milk adulteration

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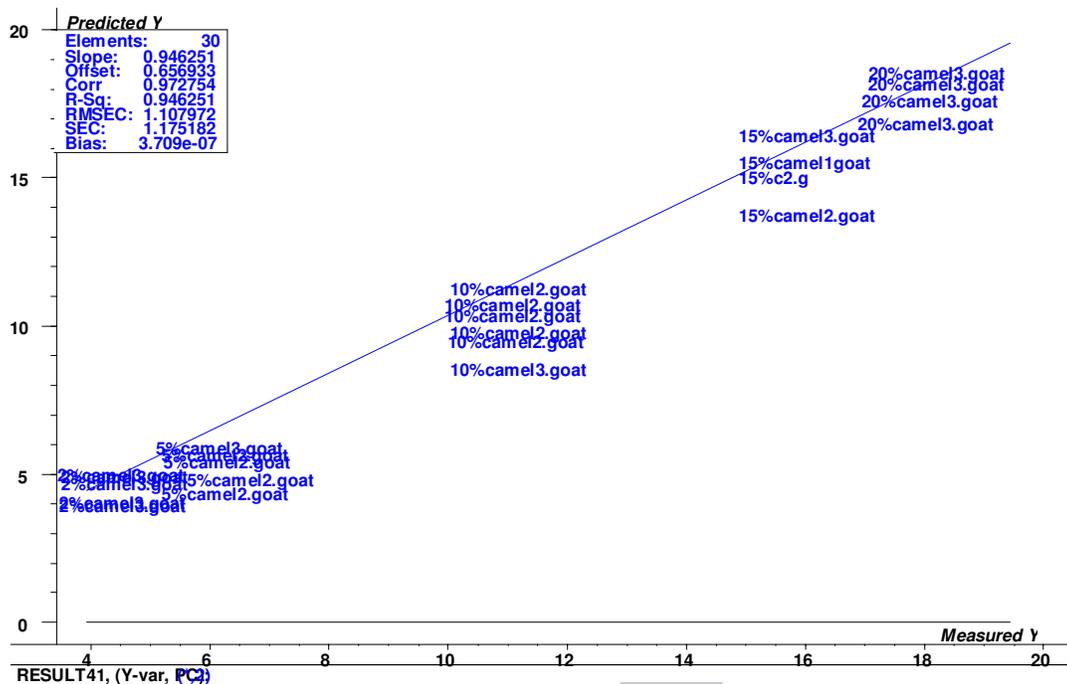
175 As Figure 3 shows a clear discrimination between the pure camel milk and with 10 %
 176 goat milk adulteration. It can be used as an identification tool to check the
 177 adulteration of camel milk with goat milk. If there is any amount of goat milk in
 178 camel milk they will occupy the space in between the pure and adulterated samples
 179 of the above PLS-DA plot. The RMSEP value for this model is 0.080 with with 0.97 R
 180 square value.

181

182 **3.3 Results of PLS regression:**

183 To quantify the amount of adulteration PLS regression model was also built by using
 184 70 % of the samples as a training set. It comprising both pure and adulterated camel
 185 milk samples 2, 5, 10, 15 and 20 % at five different percentage levels:

186



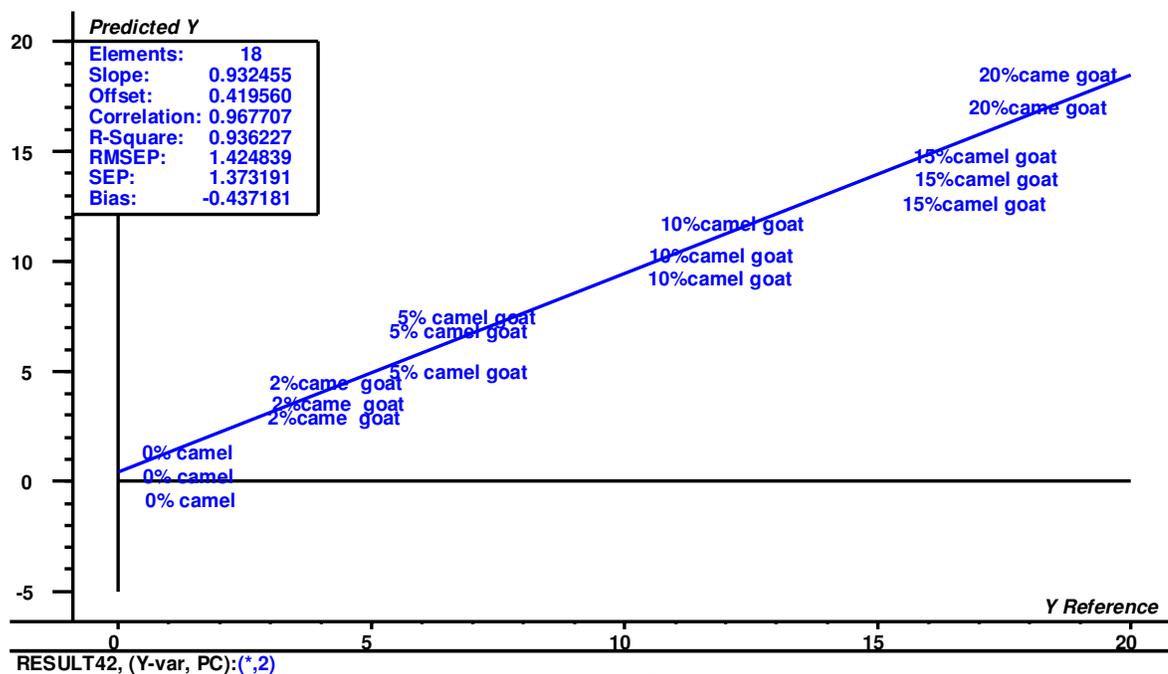
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188 Figure 4. PLS regression plot for pure and adulterated camel milk samples

189 It can be seen from Figure 4 that RMSEC = 1.10% has small value for 2 factors with
 190 $R^2 = 94\%$ and of 0.97 correlation ship.

191 To predict an independent 30% test samples set (described in the experimental
 192 section PLS calibration model was then used.

193



194

195 Figure 5 :Prediction plot for the Test set spectral data for both pure and adulterated
196 camel milk samples.

197 It can be seen from Figure 5 that the PLS calibration model is having a very good
198 prediction ability because it was applied to 30% test samples those were not used in
199 building the PLS calibration model. Again it is having minimum amount of prediction
200 error i.e. RMSEP value = 1.42% with good prediction.

201

202 4. Conclusion:

203 Newly developed NIR spectroscopic method combined with multivariate analysis
204 concludes that this is a suitable method for checking the detection and
205 quantification of camel milk adulteration with goat milk. It was further investigated
206 that PLS-DA model can be used as an identification tool while PLS calibration model
207 as a quantification. This method is simple, no need of much sample preparation and,
208 having excellent sensitivity and reproducibility.

209

210

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248

249 **Highlights**

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- Development of new NIR spectroscopic method combined with multivariate methods to detect & quantify adulteration in camel milk with goat milk
 - To build PLS regression models to quantify the amount of goat milk.
 - To build PCA model to explore the classification among various varieties of camel, cow and goat milks

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