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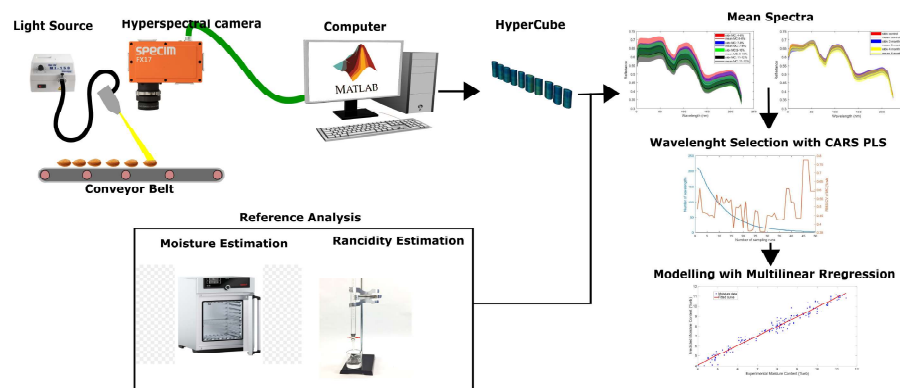
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Graphical Abstract

Rancidity and Moisture Estimation in **Shelled** Almond Kernels using NIR Hyperspectral Imaging and Chemometric Analysis*

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Highlights

Rancidity and Moisture Estimation in **Shelled Almond Kernels using NIR Hyperspectral Imaging and Chemometric Analysis**

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- Hyperspectral imaging was used to develop non-destructive method for determination of MC, FFA and PV of almond.
- Full spectrum PLSR models predicted MC, FFA and PV with R_p^2 values of 0.957, 0.970, 0.955, respectively.
- CARS-PLS was used to select the feature wavelengths for rapid and accurate quantification.
- MLR models with feature wavelengths showed R_p^2 values of 0.941, 0.903, 0.886 for MC, FFA and PV, respectively.

Rancidity and Moisture Estimation in Shelled Almond Kernels using NIR Hyperspectral Imaging and Chemometric Analysis

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Abstract

The current work focused on rapid and non-destructive determination of moisture content (MC), free fatty acids (FFA) and peroxide value (PV) in shelled almonds, using reflectance NIR-hyperspectral imaging (HSI). Sample set of 354 and 235 almond kernels were treated to vary the MC, rancidity (FFA and PV), respectively and used for model development. The sample set for each response was divided into calibration and testing set in the ratio of 70:30. Reference MC, FFA and PV were measured using wet anal-

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ysis methods and their association with the spectral data were modelled using partial least square regression. Superior models were obtained from the full-spectrum (900-1700 nm) data with R_p^2 values of 0.957, 0.970, 0.955 and $RMSEP$ values of 0.41%, 0.108%, 0.453 mEq for MC, FFA and PV, respectively. Competitive adaptive reweighted sampling method was used to select the feature wavelengths for rapid quantification. Multiple linear regression models were developed using the feature wavelengths having good predictability (R_p^2 values: 0.941, 0.903, 0.886 and $RMSEP$ values: 0.494%, 0.162%, 0.658 mEq for MC, FFA and PV, respectively). The current findings demonstrated great feasibility in industrial deployment of HSI technique for non-destructive estimation of moisture and rancidity indices in almonds and other nuts.

Keywords: Hyperspectral imaging, Moisture, Free fatty acids, Peroxide value, Non-destructive analysis

1. Introduction

Almond is the major tree nut consumed on earth, accounting nearly one third of all the tree nuts produced (inc, 2020). These are basically the fruits of almond tree (*Prunus dulcis*), grown for the seed purpose. Almonds are considered as superfood because of their rich nutritional profile containing high amount of mono-unsaturated fatty acids and polyunsaturated fatty acids, fibre, proteins, minerals and vitamins B and E (Yildirim et al., 2016). Owing to the growing health concern all around and nutritional advantages of almonds, consumers are now getting a range of option to enjoy almonds in various desserts like ice creams, chocolates, and sweets or in baked and

11 confectionery products ([Cortés et al., 2018](#)).

12 The USA, Spain, and Australia are the tabletop three exporters of al-
13 monds in the world ([inc, 2020](#)) hence carry the major responsibility of en-
14 suring the nuts qualities to be exported. This leads to need for stringent
15 regulations on accurate quality estimations which is driven by growing con-
16 cern from the consumer end and the prevailing competitive market. One of
17 the major quality concerns associated with the almond industry is rancidity
18 caused due to lipid oxidation within the nuts ([Canneddu et al., 2016](#)). There
19 could be multiple extrinsic and intrinsic factors causing rancidity such as
20 high moisture content, temperature, relative humidity, insect damage, and
21 fungal growth ([Moscetti et al., 2013](#)). However, high moisture in nuts during
22 harvest, is reportedly one of the prime reasons behind initiating hydrolytic
23 rancidity ([Canneddu et al., 2016](#); [Ghirardello et al., 2013](#)). High moisture
24 level encourages microbial growth, that produce certain enzymes responsible
25 for hydrolyzing the fat to form free fatty acids (FFAs) ([Moscetti et al., 2013](#);
26 [Wang et al., 2014](#)). Development of FFAs compromise the flavor, odor, and
27 taste of the almonds hence reduce the consumer acceptability. Along with
28 FFAs value the other most used parameter for the estimation of rancidity in
29 nuts is peroxide value (PV) which indicates the stability of the nuts against
30 oxidation ([Borompichaichartkul et al., 2013](#)).

31 Several wet lab analytical techniques are recommended for estimation of
32 fat quality ([Association of Official Agricultural Chemists & Horwitz, 1975](#);
33 [Möller, 2010](#); [ISO 17059:2019](#)) in nuts. Although these classical techniques
34 are quite reliable and accurate, have several downsides as well such as destruc-
35 tive sample preparation, involvement of hazardous chemicals, time consum-

36 ing, require large batches of samples, and limitation of accessing intrasam-
37 ple variabilities (ISO 17059:2019; Bai et al., 2018; Han et al., 2021; Car-
38 valho et al., 2019). Therefore, these classical analytical methods are losing
39 their relevance in food industries, and a complete automation and real-time
40 control during process monitoring are the need of the moment. In the re-
41 cent times, hyperspectral imaging (HSI) has emerged as a promising tool
42 for non-destructive quality estimation of food commodities (Mahesh et al.,
43 2015; Huang et al., 2014; Elmasry et al., 2012; Sricharoonratana et al., 2021;
44 Khamsopha et al., 2021).

45 HSI can operate across a wide range of wavelengths (780-2500 *nm*) de-
46 pending upon the chemical nature of the food commodity. The specific one
47 operates in the wavelength range of 900-1700 *nm* termed as shortwave in-
48 frared HSI (SWIR-HSI). SWIR-HSI acquires both spatial and spectral infor-
49 mation of the captured images and combine them to provide unique chemical
50 fingerprint to each pixel of the captured image (Elmasry et al., 2012). This
51 feature is certainly useful in dealing with the critical concern around food
52 heterogeneity (Cruz-Tirado et al., 2020). Moreover, this technique is capa-
53 ble of non-destructive evaluation of chemical composition in very short time.
54 Interestingly, some of the recent trials carried out for quality estimation of
55 nuts have shown promising outcome in favor of HSI technique (Faqeerzada
56 et al., 2020; Bilal et al., 2020; Torres et al., 2021; Moscetti et al., 2015; Jin
57 et al., 2016). HSI-based analysis produces high dimensional data set which
58 demands high computational complexity to carry out the analysis (Huang
59 et al., 2014). This situation is handled with chemometrics which allows to
60 reduce the dimensionality of the data, retain essential spectral information,

61 and classify or quantify important areas of an image (Gowen et al., 2007;
62 Amigo et al., 2008; Mishra et al., 2021).

63 Although HSI-based analysis has predominantly been applied for quality
64 estimation of agricultural produce such as fruits, vegetables, and meats, but
65 for nuts quality it is relatively underexplored (Suktanarak & Teerachaichayut,
66 2017; Teerachaichayut & Ho, 2017). Moreover, the current systems employed
67 for nuts sorting rely on random sampling that decides the market value of
68 entire batch leading to huge losses (Canneddu et al., 2016). More recently,
69 Faqeerzada et al. (2020) and Torres et al. (2021) have demonstrated applica-
70 bility of HSI on quality estimation of almond powder and shelled almonds, re-
71 spectively. However, no work in the past have addressed the non-destructive
72 estimation of both rancidity and moisture content in shelled almonds, to the
73 best of our knowledge. Hence, in the current study SWIR-HSI was used to
74 investigate the rancidity and moisture content in shelled almonds. The major
75 objectives formulated in this study were to:

- 76 • develop a full spectrum SWIR-HSI regression model for accurate pre-
77 diction of rancidity indices (i.e., FFAs and PV) and mc (%wb),
- 78 • select the feature variables for development of multispectral regression
79 model for rapid quantification,
- 80 • and visualize the spatial distribution of FFAs, PV, and mc in almond
81 kernels.

82 2. Materials and Methods

83 2.1. Almond sample preparation

84 Freshly harvested and cleaned batch of almonds of variety “Nonpareil”
85 were collected from the almond processing company Lindsay point (Victoria,
86 Australia) for the experiment. Almond samples were collected in 2019-2020
87 season. On arrival, the samples were packed in zip lock bags and stored in
88 refrigerator at 4°C until sample preparation, scanning and reference analysis.
89 To develop robust calibration model for prediction of moisture and rancidity,
90 the level of moisture content and rancidity level were varied by moisture
91 conditioning and accelerated storage, respectively.

92 Initial moisture content of the fresh almonds was determined by oven dry-
93 ing method (AOAC, 2019) and found to be $4.1\% \pm 0.15$ (wb). One kilogram
94 each batch of almond kernels was conditioned to four different moisture con-
95 tents, viz. 6, 8, 10, and 12% (wb) by adding calculated amount of **distilled**
96 **water** and thorough mixing. The conditioned almond kernels were kept in zip
97 lock low density polyethylene (LDPE) bags and stored inside moisture lock
98 compartment of refrigerator for a week in order to achieve uniform moisture
99 distribution within the kernels. The final moisture content of the kernels
100 was determined after one week of treatment by gravimetric analysis (AOAC,
101 2019). **Six almond kernels of similar moisture content were placed in a line**
102 **for scanning of each hyperspectral image. Seventy five images of each five**
103 **moisture content levels were scanned to build image library of 375 almond**
104 **images for development of the moisture regression model. During calibration**
105 **of the full-spectrum moisture prediction model, 21 outliers were removed to**
106 **get final image library of 354 almond images.**

107 Rancidity in the samples was achieved through accelerated aging of the
108 almond kernels under 75% relative humidity and 40°C temperature (Zacheo
109 et al., 1998). The almond kernels were divided into two groups, each weigh-
110 ing one kilogram were stored separately inside vacuum desiccators containing
111 saturated sodium chloride (NaCl) solutions at the base for maintaining 75%
112 relative humidity (Young, 1967). The desiccators were stored inside an en-
113 vironmental chamber maintained at 40°C and samples were removed after 3
114 months and 4 months of storage. The rancid samples were packed in zip lock
115 LDPE bags and kept in refrigerator for scanning and reference analysis. Six
116 almond kernels of same storage duration were placed in a line for scanning
117 of each hyperspectral image. Eighty images of each three storage durations
118 were scanned to build image library of 240 almond images for development
119 of the rancidity regression models. During calibration of the full-spectrum
120 rancidity prediction models, 5 and 7 outliers were removed to get final im-
121 age library of 235 and 233 almond images for FFA and PV prediction. Free
122 fatty acid content and peroxide values were considered as the indicators of
123 rancidity of the almond samples.

124 2.2. Hyperspectral image acquisition

125 The hyperspectral images of almond samples were acquired by using a
126 short-wave infrared (SWIR) hyperspectral imaging system. The system in-
127 cluded a InGaAs camera (Specim FX17, Spectral Imaging Ltd., Oulu, Fin-
128 land) with the spectral range of 900–1700 *nm* with spatial resolution of 640
129 pixels, 224 wavebands and spectral resolution of 8 *nm* full width half max-
130 imum (FWHM), one 150-watt halogen light source equipment with a fiber
131 optic line light (Illumination, Technologies Inc. USA) to illuminate the cam-

132 era’s field of view, an one-axis linear conveyor belt, and a computer (Dell,
 133 Inter(R) Core (TM) i5-2400 CPU@ 3.10 GHz, 16GB RAM) with a software
 134 called Lumo scanner (Spectral Imaging Ltd., Oulu, Finland) which controlled
 135 the system to acquire the hyperspectral images of the almond samples. Six
 136 almond kernels at a time were placed on the conveyor belt platform to be
 137 scanned with appropriate resolution, which can be maintained by moving
 138 the belt at correct speed and by maintaining correct exposure time. The
 139 halogen light source intensity was set to 60% and the exposure time was set
 140 to 12 ms during scanning. The distance between almond kernels and the
 141 lens was about 10 cm. The moving speed of conveyor belt was maintained
 142 at 15 mm/sec to obtain a hypercube of dimension 1200 × 640 × 224.

143 A dark current image (\mathbf{I}_D) and white reference image (\mathbf{I}_w) were obtained
 144 after scanning of each image by covering the camera lens and using a 99%
 145 reflectance standard (Spectralon™, SRS-99-10, Labsphere, Inc., North Sut-
 146 ton, NH, USA), respectively. Noise signals present in the raw hyperspectral
 147 images (\mathbf{I}) were corrected using equation (1) (Sun et al., 2020).

$$\mathbf{I} = \frac{\mathbf{I}_{\text{Raw}} - \mathbf{I}_D}{\mathbf{I}_w - \mathbf{I}_D} \quad (1)$$

148 2.3. Hyperspectral image segmentation and spectral data extraction

149 Segmentation was applied to the corrected images for removal of the back-
 150 ground and for extraction of the Region of Interest (ROI) in each sample
 151 using MATLAB 2019b (MathWorks Inc., Natick, MA). The difference be-
 152 tween the background and almond pixels were obtained from the HSV (hue,
 153 saturation, value) color space of the images and then a threshold value of
 154 0.45 was applied to the resultant image. A binary mask was then created

155 for each sample, assigning 0 value for the background and 1 for the pixels
156 corresponding to almonds. The mask was then applied to each hyperspectral
157 image cube before spectral data extraction. Mean spectra of each image were
158 extracted by averaging index values of each almond pixel, producing a total
159 of 224 spectral data points. Final spectral matrix for development of models
160 were obtained by concatenating the spectral data of all the almond samples.
161 In the final spectral matrix, the row represents the number of almond im-
162 ages and column represents number of wavelengths. Therefore, we obtained
163 two spectral matrices of dimension 354×224 and 235×224 for moisture and
164 rancidity value prediction, respectively.

165 *2.4. Spectral data pre-processing*

166 Prior to development of the multivariate regression model, pre-processing
167 and outlier removal techniques were performed to improve the signal to noise
168 ratio by reducing undesirable effects such as light scattering, baseline shifts,
169 random noise, and uncontrolled external factors (Bai et al., 2018). MATLAB
170 2019b (MathWorks Inc., Natick, MA) was used to pre-process all the spectral
171 data of the scanned almond images. The signal-to-noise ratio was improved
172 and useful information for models were extracted by pre-processing the spec-
173 tral data. In this study, five different pre-processing methods, viz. stan-
174 dard normal variate (SNV) and Savitzky–Golay (SG) smoothing, SG+SNV,
175 SG+first derivative, SG+second derivative were used to pre-process the raw
176 spectral data (Mishra et al., 2020). Suitable pre-processing techniques were
177 selected based on the full spectrum model performance. SNV was used to
178 reduce the baseline shift produced due to the multiplicative effects of light
179 scattering during the interaction between infrared (IR) radiation and the

180 sample particles. A constant offset term was removed by subtracting the
181 mean value of the full spectrum using SNV, which brought all spectra to
182 the same scale by subsequent division by the standard deviation of the full
183 spectrum (Grisanti et al., 2018). Moreover, Savitzky–Golay filter was applied
184 to smoothing the spectrum in addition with 1st and 2nd derivative, which
185 enhanced the difference between the reflectance values.

186 *2.5. Reference analysis*

187 Each sample set of almonds after hyperspectral image acquisition were
188 ground in a laboratory mixer and packed in zip lock LDPE bags, individually.
189 The ground almond samples were used in the measurement of moisture and
190 rancidity analysis as described below.

191 *2.5.1. Moisture analysis*

192 Ground almond samples were analyzed for moisture content using stan-
193 dard gravimetric methods described by AOAC (2019). Approximately five
194 grams of ground almond for each sample **in duplicates were** placed in pre-
195 weighed aluminium moisture box and put into a drying oven (DHG-9101-
196 2SA; Shanghai Sanfa Scientific Instruments Co., Ltd, Shanghai, China) main-
197 tained at 130°C for 3 hours. After drying, the moisture boxes were imme-
198 diately cooled in a vacuum desiccator containing allochronic silica gel for 1
199 hour. The weight of the ground almond samples was recorded before and after
200 drying using weighing balance (company and sensitivity), and the moisture
201 content was determined by the mass loss after drying. Wet basis moisture
202 content was calculated using the following equation:

$$\text{Moisture content}(\%wb) = \frac{W_2 - W_1}{W_3 - W_1} \times 100 \quad (2)$$

203 where W_1 is the weight of empty moisture box, W_2 and W_3 are weights of
204 moisture box with almond samples before and after drying, respectively. All
205 weights are in grams.

206 2.5.2. Rancidity analysis

207 Free fatty acid (FFA) content and peroxide values (PV) were used as
208 rancidity index for almond samples in present study. Almond oil from each
209 set of samples were extracted using a hydraulic press and the extracted oils
210 were immediately analyzed for the rancidity indices. For FFA analysis, ap-
211 proximately 2 g of oil (W) was extracted from almond samples **in duplicates**
212 **were** placed into a 100 mL Erlenmeyer flask and mixed with 10 mL ether, 10
213 mL 95% ethanol, 2 to 3 drops of 1% phenolphthalein indicator. The mixture
214 was titrated with 0.005 mol/L sodium hydroxide ethanol solution until the
215 mixture turns pink for at least 1 min. The volume of the titrant consumed
216 was recorded as V . FFA content was calculated using the following equation,
217 where C is the calibrated concentration of sodium hydroxide ethanol solution
218 and 282 is oleic acid molar mass in g/mol:

$$FFA(\%) = \frac{V \times C \times 282}{1000 \times W} \times 100 \quad (3)$$

219 Peroxide value was estimated by using the titration method provided in
220 AOAC Official Method 965.33 (AOAC, 2019) with a slight modification to be
221 used for microtitration. Approximately, 2 g of oil sample (W) **in duplicates**
222 **were** placed into a 100 mL Erlenmeyer flask and mixed with 12 mL acetic
223 acid–chloroform (3:2) solution, and 0.5 mL saturated potassium iodide (KI)
224 solution. The solution was allowed stand with occasional swirling for 1 min
225 and then 12 mL distilled water was added. After 1 min, the solution was

226 slowly titrated with 0.01-N sodium thiosulfate ($Na_2S_2O_3$). Continue titrat-
 227 ing until the yellow color goes away. 0.5 mL of 1% soluble starch indicator
 228 was then added to change the color from light yellow to purple. The solution
 229 was again titrated with sodium thiosulfate ($Na_2S_2O_3$), dropwise until the
 230 purple color just disappears. The *PV* as mEq of oleic acid per kg of oil was
 231 calculated according to following equation:

$$PV = \frac{(V \times C \times 1000)}{W} \quad (4)$$

232 where V is the volume of $Na_2S_2O_3$ solution in mL, C is the concentration of
 233 the $Na_2S_2O_3$ solution in mol/L, and W is the weight of the oil sample taken
 234 in g.

235 2.6. Full spectrum calibration model development

236 PLSR model was used to establish the relationship between the spectral
 237 data and the response, viz. moisture, FFA and PV. For each parameter, SG-
 238 second derivative pre-processed spectral data (\mathbf{X}) and the reference response
 239 values (\mathbf{Y}) were used for the regression model development. The sample set
 240 for each response was divided randomly into two categories, viz. calibration
 241 set, and testing set. The raw spectra were projected onto several latent
 242 variables (LVs) for simplifying the association of spectra and the attribute
 243 matrixes. The LVs maximize the covariance between the two matrices of the
 244 original data and significantly reduce the original spectral dimension. The
 245 ideal number of LVs is decided by cross validation when the root mean square
 246 error of cross-validation ($RMSE_{CV}$) reaches a minimum. In this study, 10-
 247 fold cross-validation was conducted and the developed model was tested by
 248 the prediction set. The association between spectra matrix \mathbf{X} ($n \times m$) and

249 quality attribute \mathbf{Y} ($n \times 1$) can be predicted using equations (5) to (7) (Wold
250 et al., 2001; Guebel & Torres, 2013).

$$251 \quad \mathbf{X}_{(n;p)} = \widehat{\mathbf{X}}_{(n;p)} + \mathbf{E}_{(n;p)} = \mathbf{T}_{(n;k)} \mathbf{P}_{(n;p)}^T + \mathbf{E}_{(n;p)} \quad (5)$$

$$\mathbf{Y}_{(n;m)} = \widehat{\mathbf{Y}}_{(n;m)} + \mathbf{F}_{(n;m)} = \mathbf{U}_{(n;k)} \mathbf{Q}_{(k;m)}^T + \mathbf{F}_{(n;m)} \quad (6)$$

$$\mathbf{U}_{(n;k)} = \mathbf{T}_{(n;k)} \mathbf{C}_{(k;k)} \quad (7)$$

252 Where $\widehat{\mathbf{X}}$ and $\widehat{\mathbf{Y}}$ give the PLS estimations of the matrices \mathbf{X} and \mathbf{Y} respec-
253 tively, \mathbf{E} and \mathbf{F} are the residual error matrices, \mathbf{T} and \mathbf{U} are the score ma-
254 trices, \mathbf{P} and \mathbf{Q} are the loading matrices. After this calculations the matrix
255 B_{PLS} of regression coeffiennts can be calaculated with crefeqbeta3,eqbeta4

$$256 \quad \mathbf{B}_{PLS} = \mathbf{W}(\mathbf{P}^T \mathbf{W})^{-1} \mathbf{C} \mathbf{Q}^T \quad (8)$$

$$\widehat{\mathbf{Y}} = \mathbf{T} \mathbf{C} \mathbf{Q}^T = \mathbf{X} \mathbf{B}_{PLS} \quad (9)$$

257 Developed PLSR models were then tested using the mean spectra of the
258 testing data set. The predictability of the developed models was evaluated
259 by using the coefficient of determination (R^2) and root mean square error
260 ($RMSE$) using the following equations:

$$R^2 = 1 - \frac{\sum_{i=1}^n (Y_i - \widehat{Y})^2}{\sum_{i=1}^n (Y_i - \bar{Y})^2} \quad (10)$$

$$261 \quad RMSE = \sqrt{\sum_{i=1}^n \frac{(Y_i - \widehat{Y})^2}{n}} \quad (11)$$

262 The co-efficients of determination of calibration (R_c^2), cross-validation
263 (R_{cv}^2) and prediction (R_p^2) and the corresponding root mean square error of

264 calibration ($RMSE_C$), root mean square of cross validation ($RMSE_{CV}$), and
 265 prediction ($RMSE_P$) were used for evaluating developed PLS models. Best
 266 fit models have high R_c^2 , R_{cv}^2 , R_p^2 and low $RMSE_C$, $RMSE_{CV}$ and $RMSE_P$
 267 together with slight differences among them. It always expects R^2 to go to 1
 268 and $RMSE$ to go to 0. Furthermore, the robustness of the model was evalu-
 269 ated by calculating the residual predictive deviation (RPD), which is the ratio
 270 between standard deviation (SD) of the population's reference value and the
 271 standard error of prediction of the cross-validated data set (Equation (12)).
 272 RPD provides standardization of prediction accuracy of the model (Mishra
 273 et al., 2018). Williams & Sobering (1996) suggested that RPD values of
 274 2.4–3.0 are poor; > 3 indicate that the model is appropriate for screening
 275 and is robust.

$$RPD = \frac{SD}{RMSE} \quad (12)$$

276 2.7. Feature wavelength selection

277 The acquired hyperspectral images contained 224 number of wavelengths,
 278 which is of high dimension and involves multi-collinearity resulting in increase
 279 in the speed of the acquisition. The high dimensional hyperspectral data is
 280 not suitable for commercial application, where the estimation speed should
 281 match with the production capacity. To improve the calculation speed and
 282 prediction accuracy, some sensitive wavelengths were selected from the full
 283 spectral matrices to compress hyperspectral data. These feature wavelengths
 284 must carry most of the useful information which could perfectly predict the
 285 moisture content and rancidity indices of the almond samples. Competitive
 286 adaptive reweighted sampling (CARS) technique was used for selection of key
 287 wavelengths. To perform CARS, MATLAB 2019b (MathWorks Inc., Natick,

288 MA) was used.

289 CARS is a useful feature selection technique, works through selecting N
290 subsets of independent variables by N sampling runs in an iterative man-
291 ner and develop a robust PLSR (Tao et al., 2019). It chooses the subset
292 with the lowest root-mean-square error of cross validation ($RMSE_{CV}$) as
293 the optimal subset. In this manuscript, the optimal wavelengths imply the
294 feature wavelengths with large absolute coefficients in the developed PLSR
295 model. CARS have **five** basic steps involved during each sample analysis:
296 (1) divide the sample set into Monte Carlo technique to the sample set, (2)
297 build PLS model at each sampling run (3) utilize exponentially decreasing
298 function (EDF) for wavelength selection; (4) adopt reweighted sampling for
299 wavelength selection in a competitive manner; and (5) conduct cross valida-
300 tion to evaluate the prediction capability of the selected wavelengths. In this
301 study, 10-fold cross validation was used to evaluate the feature wavelengths,
302 and the number (N) of Monte Carlo sampling runs was set to be 50. The pro-
303 cess was repeated The PLSR model was re-developed using the reflectance
304 value of feature wavelengths to predict the moisture and rancidity indices
305 of the almond samples, rapidly. Then the developed model was tested by
306 applying on the external testing tests.

307 *2.8. Development of multiple linear regression (MLR) model using feature* 308 *wavelengths*

309 MLR models were established for MC, FFA and PV using the selected
310 wavelengths from CARS, without any pretreatment in order to simplify pre-
311 diction process and to improve calculation speed. Testing data set were
312 used for development of calibration model and were validated using 10-fold

313 cross-validation process. Testing data sets were used to predict the MC,
314 FFA and PV of almond kernels. R_c^2 , R_{cv}^2 , R_p^2 and the corresponding $RMSE_C$,
315 $RMSE_{CV}$, and $RMSE_P$ values were used for evaluating developed MLR mod-
316 els.

317 *2.9. Classification of almonds according to rancidity*

318 Support vector machine classification (SVM) were performed to classify
319 almonds based on rancidity parameters, viz. FFA value and PV using the
320 feature wavelengths. Two class classification was done considering class A as
321 fresh having FFA value $<1\%$ and $PV < 5$ Meq/kg and class B as rancid having
322 FFA value $>1\%$ and $PV > 5$ Meq/kg. SVM is a machine learning technique
323 used to handle nonlinear input spaces. SVM constructs a hyperplane in
324 multidimensional space to separate different classes. SVM generates optimal
325 hyperplane in an iterative manner, which is used to minimise an error. The
326 core idea of SVM is to find a maximum marginal hyperplane (MMH) that best
327 divides the dataset into classes (Cruz-Tirado et al., 2021). The calibration
328 set was used to build the model with 10-fold internal cross-validation and
329 prediction was performed using 30% of remaining test samples. R_c^2 , R_{cv}^2 , R_p^2
330 and the corresponding $RMSE_C$, $RMSE_{CV}$, and $RMSE_P$ values were used for
331 evaluating developed MLR models.

332 *2.10. Visualization of MC and rancidity distribution*

333 Visualization of the moisture and rancidity distribution in a batch of
334 sample, the calibration model was first built using the mean spectrum of
335 all pixels of each seed and the reference MC obtained by the gravimetric
336 method as described in the previous sections. Then the MC value, FFA and

337 PV content of each pixel were predicted by the developed PLS model. MC
338 and rancidity distribution maps were created with a pseudo color bar and the
339 spatial distribution of MC, FFA and PV can be assessed by observing color
340 change in a distribution map. Pseudo color maps for quality distribution
341 were developed using MATLAB 2019b (MathWorks Inc., Natick, MA).

342 **3. Results and Discussion**

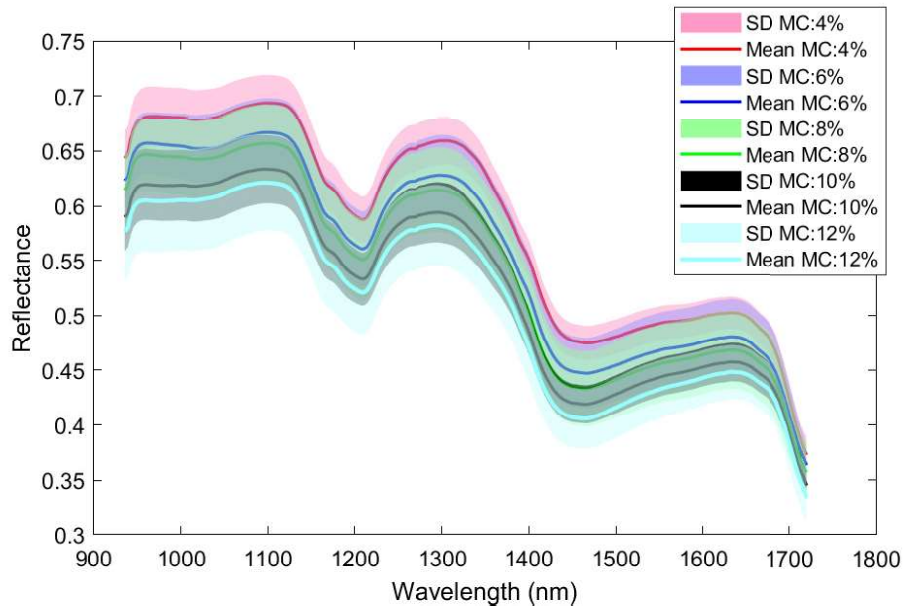
343 *3.1. Spectral features*

344 The reflectance spectral library obtained from almond samples with **five**
345 **different MC levels, viz. 4, 6, 8, 10, 12% (wb) and three different rancidity**
346 **levels, viz. control, stored for 3 months and four months** are shown in fig-
347 ure **1a** and figure **1b**, respectively. The most <https://www.overleaf.com/project/615b465d52462e38a>
348 wavebands existing in the NIR region are the strong overtone and combina-
349 tion absorptions of hydrogen containing bonds O–H (found in water), C–H
350 (found in carbohydrates & oil), and N–H (found in protein). The reflectance
351 values gradually decreased with the increase in MC throughout the spectra
352 (Figure **1a**) (Xu et al., 2019). **In figure 1a, the whole spectra are found**
353 **to be varied with varying moisture content as water greatly absorbed NIR**
354 **radiation and O-H bond dominated the NIR spectra of kernels. The lumi-**
355 **nous energy from the NIR electromagnetic waves is transferred to the elastic**
356 **potential energy (stored in the form of bending O–H bonds) and chemical po-**
357 **tential energy (stored in the form of stretching O–H bonds). Due to increase**
358 **moisture content in the kernel, more energy from electromagnetic waves is**
359 **absorbed, reflecting less energy back into the lens of the hyperspectral cam-**
360 **era (Deng et al., 2015). Thus, it interferes with the reflectance characteristics**

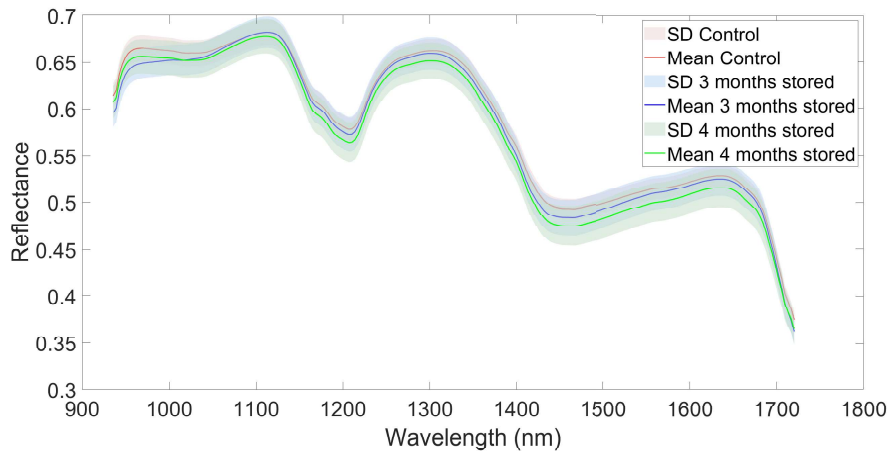
361 of other molecules present in the kernel. The absorption peaks at 960 *nm*
362 and valleys at 1450 *nm* are associated with the second and first overtone of
363 O–H stretches in water, respectively (Zhang & Guo, 2020). The reflectance
364 peaks or valleys of rancid almond spectra increased at specific wavebands,
365 viz. 1210, 1450, and 1638 *nm* with increase in the rancidity level (figure 1b),
366 however all other peaks were not showing any relationship among them. The
367 absorption bands at 1200-1250 *nm* are related to C-H stretch second over-
368 tone and the spectral region between 1600 and 1650 *nm* is related to the
369 first overtone of C–H vibration, influenced by the presence of CH₃ bonds
370 of fats present in the almond kernels (Cruz-Tirado et al., 2020). The ab-
371 sorption peak at 920-950 *nm* probably related to the combination effect of
372 third overtone of band C-H and fat (Sun et al., 2020). The change in the
373 spectral features in the rancid almond samples were observed might be due
374 to oxidation of the complex fat into free fatty acids and peroxides because
375 of storing the almonds in a high humidity and temperature condition (Hong
376 et al., 2018). Consequently, more lipid degradation reported as the storage
377 duration increased from 3 months to 4 months. The variation in the absorp-
378 tion peaks of rancid almonds at 960, 1150 and valleys at 1450 *nm* might be
379 due to the O-H stretching, indicating that there was a reduction in moisture
380 content during storage. Similar spectral variation was also found by Carvalho
381 et al. (2019) for macadamia nuts.

382 3.2. Full-spectrum PLS models of MC

383 Final moisture content obtained after conditioning were vary from $4.52 \pm 0.45\%$
384 (wb) for control samples to $11.44 \pm 0.5\%$ for samples conditioned to 12%(wb)
385 (table 1). Based on the all the pre-processed spectral data extracted from



(a)



(b)

Figure 1: Mean reflectance spectra with standard deviation obtained from almond kernels of (a) treated moisture and (b) stored in accelerated conditions (N.B: SD=Standard deviation; MC=moisture content in % wet basis)

386 the hyperspectral images of almond kernels and their corresponding reference
387 MC values, the calibration models with the whole spectral wavelengths were
388 established using PLSR algorithms. table 2 shows the calibration, cross-
389 validation, and prediction outcome of the above two established models.
390 Calibration plot obtained for training MC is shown in figure 2a. The model
391 performances were given in terms of R_c^2 , R_{cv}^2 , R_v^2 and corresponding $RMSE_C$,
392 $RMSE_{CV}$, and $RMSE_P$ values. SG+1st derivative was found to be the best
393 pre-processing techniques for prediction of MC, with seven LVs, R_p value
394 of 0.957 and $RMSE_P$ value of 1.702%. The model developed using SG+1st
395 derivative also had high RPD value of 5.07, showing the robustness of the
396 model. All other pre-processing techniques showed poor performance having
397 low RPD values as compared to the raw spectra. **The reason behind poor per-**
398 **formance on application of SNV pre-processing technique alone or together**
399 **with SG filter might not properly normalise the spectra since within-class**
400 **mean spectrum might be inappropriate in the particular data set. Thus,**
401 **the applied pre-processing might cause interaction of the unimportant infor-**
402 **mation, resulting non-orthogonal nature of the data.** The prediction result
403 obtained from the spectral data indicated that SG filter with combination of
404 derivation pre-processing techniques improved the PLS model predictive ac-
405 curacy, which might be due to smoothing of the spectra and enhancement of
406 the difference between them. The prediction error for natural moisture con-
407 tent was always much below 0.3% for a range of 4-5%, whereas for laboratory
408 treated almond kernels, the prediction error was 0.6% for a moisture range
409 of 11-12% and thus, the average prediction error was found to be 0.410. The
410 capability of HSI for moisture prediction is consistent with previous studies

411 of (Xu et al., 2019) and (Sun et al., 2020) for who reported higher prediction
 412 accuracy for cucumber seeds and peanuts with R_p^2 values of 0.917 and 0.943,
 413 respectively.

Table 1: Statistics of moisture content and rancidity indices of various sample sets

Sample set	No. of samples	Min	Max	Mean	SEM	CV
MC (% wb)						
Total	354	4.00	11.44	7.72±2.01	0.10	0.26
Calibration	250	4.00	11.44	7.66±1.99	0.12	0.25
Testing (Prediction)	104	4.40	11.44	7.89±2.05	0.20	0.27
FFA (%)						
Total	235	0.20	1.62	1.23±0.51	0.03	0.42
Calibration	165	0.20	1.62	1.26±0.53	0.04	0.44
Testing (Prediction)	70	0.20	1.61	1.14±0.48	0.05	0.37
PV (Meq OAA)						
Total	233	1.91	7.62	3.43±2.01	0.13	0.58
Calibration	165	1.91	7.62	3.62±2.00	0.15	0.59
Testing (Prediction)	68	1.91	6.99	2.97±2.03	0.24	0.58

N.B. Mean±SD indicates the average value with standard deviation among the sample set; SEM=standard error of mean; CV=Co-efficient of variation

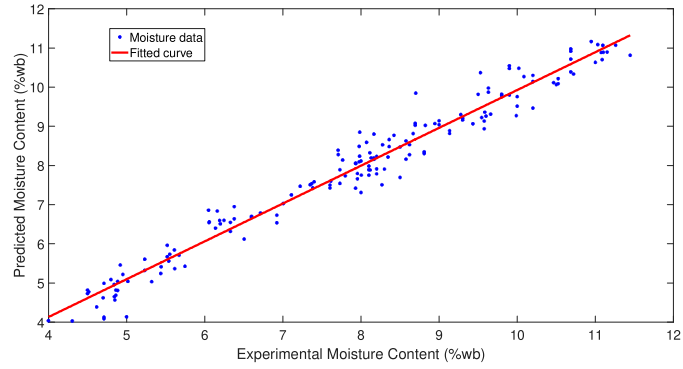
414 3.3. Full-spectrum PLS models of rancidity indices

415 Rancidity of the stored samples were calibrated according to the FFA
 416 and PV values of each storage period. FFA values of the stored samples were
 417 increased drastically after 3 months of storage, but peroxide values were

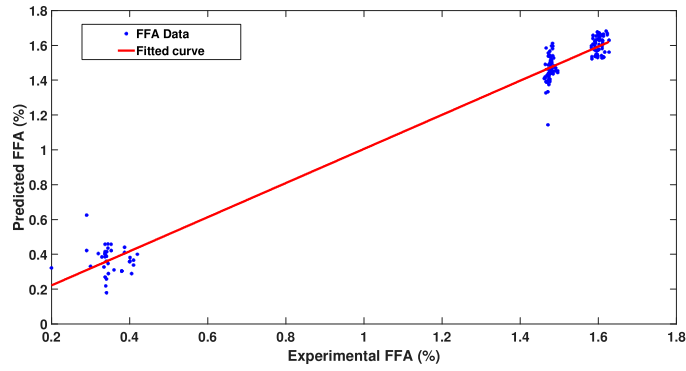
Table 2: PLS prediction results of almond MC and rancidity with different pretreatments

	Pre-Processing Methods	LVs	Calibration		Cross-validation		Prediction		RPD
			R_c^2	$RMSE_C$	R_{cv}^2	$RMSE_{CV}$	R_p^2	$RMSE_P$	
Moisture content (%wb)	Raw	9	0.967	0.375	0.944	0.464	0.943	0.441	4.550
	SNV	9	0.961	0.384	0.947	0.443	0.959	0.432	4.070
	SG	10	0.917	0.567	0.900	0.607	0.859	0.778	3.310
	SG_SNV	8	0.900	0.627	0.881	0.653	0.884	0.698	3.150
	SG_1st derivative	7	0.973	0.328	0.966	0.366	0.957	0.410	5.070
	SG_2nd derivative	7	0.970	0.347	0.954	0.432	0.946	0.451	4.330
FFA (%)	Raw	10	0.958	0.101	0.945	0.108	0.931	0.123	4.020
	SNV	8	0.970	0.091	0.954	0.111	0.927	0.124	4.93
	SG	9	0.975	0.082	0.972	0.083	0.970	0.053	5.600
	SG_SNV	7	0.983	0.066	0.977	0.074	0.958	0.108	5.840
	SG_1st derivative	7	0.971	0.087	0.959	0.102	0.948	0.113	5.060
	SG_2nd derivative	8	0.976	0.079	0.958	0.015	0.943	0.122	5.050
PV (Meq OAA)	Raw	10	0.943	0.463	0.927	0.508	0.934	0.535	3.940
	SNV	9	0.950	0.447	0.925	0.543	0.928	0.533	3.740
	SG	8	0.974	0.313	0.969	0.337	0.941	0.410	4.870
	SG_SNV	8	0.975	0.349	0.962	0.382	0.955	0.453	5.630
	SG_1st derivative	9	0.952	0.453	0.934	0.519	0.920	0.506	4.410
	SG_2nd derivative	8	0.973	0.328	0.949	0.439	0.907	0.615	3.840

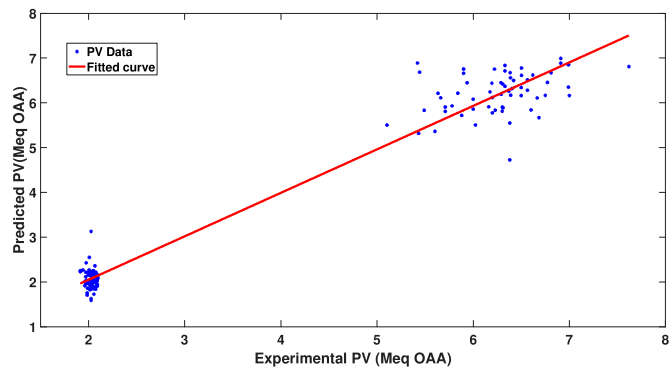
LV=latent variable; R_c^2 , R_{cv}^2 and R_p^2 =regression co-efficients of calibration, cross-validation and prediction, respectively; $RMSE_c$, $RMSE_{cv}$, $RMSE_p$ =root mean square error of calibration, cross-validation and prediction, respectively; RPD=residual predictive deviation



(a)



(b)



(c)

Figure 2: Experimental vs predicted results obtained from the full spectrum PLSR models for (a) MC, (b) FFA and (c) PV

418 not increased significantly even after 3 months of storage; however, after 4
419 months of storage PV content was also increased above 5 Meq OAA. FFA
420 value obtained after 3 and 4 months storage was about $1.47\pm 0.006\%$ and
421 $1.60\pm 0.010\%$, respectively. Moreover, the PV of the almonds were found
422 to be $2.03\pm 0.03\%$ and $6.27\pm 0.43\%$, respectively for 3 and 4 months stor-
423 age samples. The results for FFA and PV model development using HSI
424 in intact almond kernels are reported in table 2. It was found that all the
425 pretreatment techniques improved the prediction capability of the calibration
426 models by removing nonlinear spectroscopic artefacts, viz. drifting baselines,
427 scattering and noise involved in the spectra. RPD values of all the models
428 using different pre-processing techniques are found to be greater than the
429 raw spectral model. The SG smoothing combined with SNV pre-treatment
430 gave the best model performance for both FFA and PV due to removal of
431 high-frequency noise during the normalisation process. Higher accuracy was
432 found using SNV pretreatment might be due to correction of baseline shift in
433 the dataset resulting from light scattering. Calibration plot showing exper-
434 imental rancidity indices vs predicted indices, using the best pre-processing
435 technique obtained are shown in figure 2b and figure 2c. Results reported in
436 figure 2b and figure 2c should be considered with caution, given the lack of
437 samples with intermediary values of reference, as this could probably caused
438 a biased prediction model with two major regions of concentration of the
439 reference data. The best full spectrum model for FFA resulted R^2 values of
440 0.983, 0.977, and 0.958 for the calibration, cross-validation, and prediction
441 datasets, respectively. The optimized calibration model for PV resulted R^2
442 values of 0.975, 0.962, and 0.955 for the calibration, cross-validation, and

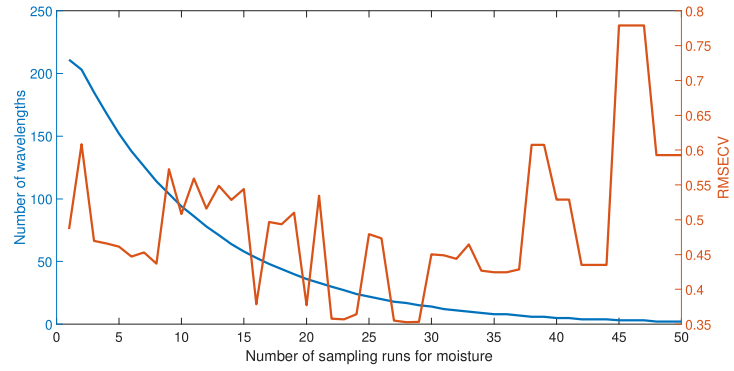
443 prediction datasets, respectively. The corresponding $RMSE_P$ and RPD val-
444 ues for both FFA and PV were found to be 0.108% and 0.453 Meq OAA, and
445 5.84 and 5.63, respectively indicating excellent performance of the developed
446 PLS models. Previous research on determination of lipid oxidation using HSI
447 also described its higher predictive capability. Han et al. (2021) reported ap-
448 plication of HSI coupled with convolutional neural network (CNN) for PV
449 estimation in canarium nut having highest overall accuracy of 93.48%. Xia
450 et al. (2016) and Ma et al. (2015) also reported superior prediction and clas-
451 sification efficiency of multispectral imaging for butter cookies and sunflower
452 seeds, respectively. They used selective wavelengths for determination of
453 rancidity indices and found peroxide value prediction accuracy of 97.1% and
454 rancid kernels classification accuracy of $> 97\%$, respectively.

455 3.4. Feature wavelength selection

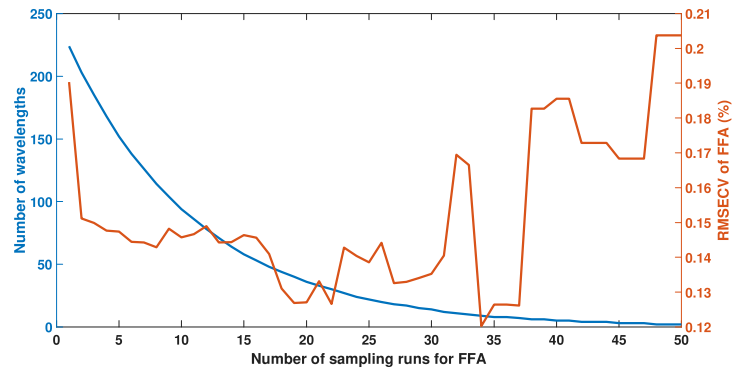
456 The selected wavelengths were compared with the β -coefficient values
457 extracted from the full spectrum PLS models. Figure 3 shows the feature
458 wavelength selection for curves for MC, FFA and PV. The blue curves in
459 Figure 3 indicating the decrease in the number of wavelength variables as
460 the number of sampling runs increases. The declining trend was found to
461 be gradually slowing down, which showed the fast and the refined selection
462 of wavelengths. The red curves in Figure 3 show the variation of $RMSE_{CV}$
463 values with the number of sampling. As the elimination process of redundant
464 wavelengths was increased, $RMSE_{CV}$ values were first reduced moderately
465 and then gradually increased with the elimination of feature wavelengths.
466 The optimum wavelengths were chosen based on the minimum $RMSE_{CV}$
467 values. For moisture calibration set, the $RMSE_{CV}$ value was lowest at the

468 29th sampling and the corresponding number of feature wavelengths were
469 7. Similarly, for rancid almond sample set, least $RMSE_{CV}$ value was found
470 at the 34th and 42nd sampling for FFA and PV, respectively and the opti-
471 mum number of feature wavelengths selected were 5 for both the parameters.
472 The feature wavelengths were selected from the optimal subset based on the
473 highest weight coefficients applying “survival of the fittest” principle. The
474 specific effective wavelengths selected by CARS technique were listed in ta-
475 ble 3. The selection of the most important wavelengths was also compared
476 with the β -coefficients of these PLS regression prediction models. Only those
477 wavelengths with large β -coefficient values are useful for the prediction and
478 are shown in figure S1.

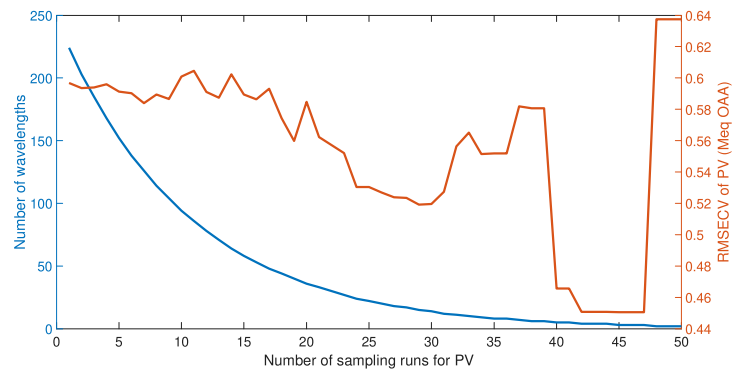
479 For moisture prediction, the feature wavelengths were 970, 1001, 1154,
480 1312, 1350, 1437, 1670 nm . Previous research on MC prediction for corn
481 kernels described the major wavelengths at 970 nm , 1150 and 1450 nm , re-
482 lated to the 2nd, 3rd and 1st overtone of O-H stretching, respectively (Zhang
483 & Guo, 2020). Our results were also comparable with the study of Liu et al.
484 (2014), who reported 1178, 1299, 1363, 1389, 1643, 1788, 1941, 2213, 2439
485 nm were the important wavelengths selected for MC determination in pork
486 samples, and their regression model showed MC prediction with a test R^2
487 of 0.920. The strongest feature wavelengths for prediction of FFA and PV
488 were 945, 994, 1105, 1382, 1527 nm and 945, 1025, 1393, 1488, 1695 nm ,
489 respectively. From β coefficient graph of FFA and PV, small peaks were
490 found at 1210 nm , which may also relate to the change in C-H bond present
491 in the almond lipids. The results were like the findings of Caporaso et al.
492 (2018), who reported the important peaks for lipid content prediction lie at



(a)



(b)



(c)

Figure 3: The result of variable selection by CARS for (a) MC (b) FFA and (b) PV in almond kernels (N.B: RMSECV=Root mean square error of cross-validation; MC=Moisture content; FFA=free fatty acids; PV=peroxide value; Meq OAA=Milli equivalent oleic acid)

493 1210, 1360, 1700 nm , mostly arise from the first and second overtone of C-H
494 and stretching of the -CH₂ groups. [Khodabux et al. \(2007\)](#) also reported
495 the absorption peaks around 1160 nm was attributed to -HC=CH- bonds.
496 Feature wavelength of 1488, 1695 nm for PV prediction might be arise due
497 to the secondary vibration resulted due to stretching of O-H bond present in
498 hydroperoxides in the samples.

499 3.5. MLR model development, validation, and testing

500 Table 3 shows the performance of MLR models built for MC, FFA and
501 PV using the selected wavelengths. The use of lower number of bands led to
502 a satisfactory performance compared to the PLS regression model reported
503 in the previous section for moisture using the full spectra range. In fact, the
504 moisture model had a $RMSE_C$ of 0.452% and $RMSE_P$ of 0.494%, which is
505 comparable to the full spectrum PLS regression model, having $RMSE_P$ value
506 of 0.410. Similarly, $RMSE_P$ values of 0.162% and 0.658 Meq OAA were ob-
507 tained from MLR model of FFA and PV, which are bit high as compared to
508 the full spectrum model, still it is acceptable provided the quick prediction
509 ability in a less complex manner. It should be noted that this prediction is
510 made on the reflectance data with no spectral pre-treatment, which result
511 into a larger prediction error when using multispectral camera arrangement
512 with few feature wavelengths. Considering the advantage of low-cost multi-
513 spectral use and the rapid computational capacity required, the seven-band
514 model for MC and five-band model for rancidity indices could be still con-
515 sidered as acceptable for practical applications for rapid screening of higher
516 or lower moisture almond kernels and rancid or fresh kernel during storage
517 and processing.

Table 3: Feature wavelengths selected by CARS PLS for MC and rancidity indices of almonds

Quality Parameters	LVs used	Selected wavelengths (nm)	Calibration		Cross-validation		Prediction	
			R_c^2	$RMSE_C$	R_{cv}^2	$RMSE_{CV}$	R_p^2	$RMSE_P$
MC	8	970, 1001, 1154, 1312, 1350, 1437, 1670	0.95	0.452	0.93	0.518	0.941	0.494
FFA	7	945, 994, 1105, 1382, 1527	0.92	0.146	0.9	0.165	0.903	0.162
PV	7	945, 1025, 1393, 1488, 1695	0.9	0.666	0.89	0.675	0.886	0.658

R_c^2 , R_{cv}^2 , and R_p^2 =regression co-efficients of calibration, cross-validation and prediction, respectively; $RMSE_C$, $RMSE_{cv}$, $RMSE_P$ =root mean square error of calibration, cross-validation and prediction, respectively

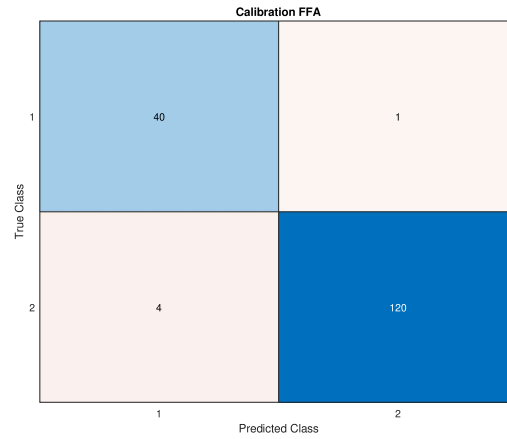
518 3.6. Classification of almonds according to rancidity

519 Figure 4 shows the confusion matrix of the two-class SVM classification
520 model for FFA and PV value indicating whether the almonds were fresh
521 or rancid using the feature wavelengths. SVM calibration models showed
522 accuracy of 97% and 97.6% in classifying the rancid almonds from the fresh
523 kernels based on FFA and PV value, respectively. The classification results
524 of FFA model showed 4 out of 44 fresh samples were classified incorrectly and
525 1 out of 121 rancid samples is classified as fresh. The the validation results
526 of PV model showed 2 out of 109 samples were classified incorrectly and 2
527 out of 56 rancid samples is classified as fresh. According to PV value the
528 classification results were different because even after 3 months of storage,
529 the PV value was not significantly increased as described earlier. Only the
530 samples stored for 4 months were considered as rancid, in which the PV
531 value exceeded 5 Meq OAA/kg. Samples classified as rancid according to
532 FFA value and PV were considered as the almonds affected by oxidative
533 rancidity and hydrolytic rancidity, respectively(Franklin & Mitchell, 2019).
534 The goodness of fit of the developed models can be demonstrated using R_v^2
535 values of 0.96 and 0.98 for FFA and PV, respectively for the external test

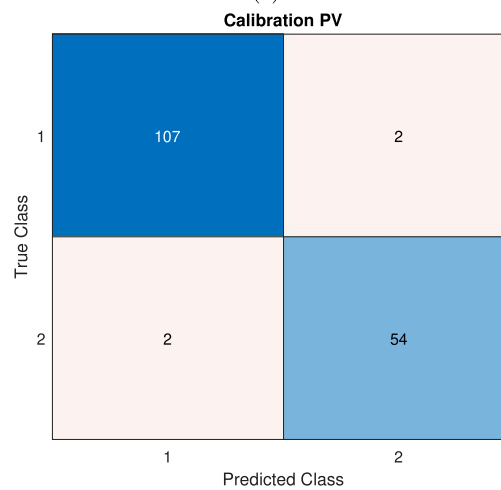
536 set. However, these models could suffer some biased overfitting caused by
537 the lack of samples in intermediate regions of the results. The classification
538 results demonstrate the potential of the multispectral imaging system with
539 the developed models to identify rancid almonds, which is very useful for
540 the automation of quality control system in nuts industries. The results
541 obtained from this work were lie in line with the previous work using portable
542 NIR spectrometer by Cruz-Tirado et al. (2021) for classification of fresh and
543 stale eggs. They developed classification models using PLS-DA and SVM to
544 discriminate between fresh (Haugh Unit > 60) and stale (Haugh Unit < 60)
545 based on Haugh Unit as a freshness indicator eggs and achieved an accuracy
546 higher than 75% of correct classification of fresh samples.

547 3.7. Visualization of the moisture distribution

548 Figure 5 shows the moisture and rancidity distribution pseudo color map
549 generated by applying the MLR models to each pixel of the multispectral im-
550 age obtained by combining the images at feature wavelengths. The purpose
551 of showing these charts is to provide the readers an idea about intra-kernel
552 distribution of the moisture and rancidity indices. It should be noted that
553 the values assigned for the specific pixels are not representing the exact con-
554 centration of moisture, FFA, and PV; instead, it is showing their relative
555 distribution across individual almond kernels. In figure 5a, the average kernel
556 moisture contents of the almonds presented in different columns were 4, 6,
557 8 and 10% (wb), respectively from left to right. As the moisture content
558 increases, the distribution of intra-kernel moisture can be easily seen. Noise
559 involved in the moisture distribution pattern of first two columns of figure 5a
560 could be due to several contributing factors such as inconsistent intra-cellular



(a)



(b)

Figure 4: SVM classification results of almond rancidity based on FFA and PV: (a) confusion matrix of classification based on FFA (%) and (b) confusion matrix of classification based on PV (Meq OAA); class '1' and '2' in the figure refers to 'fresh' and 'rancid' almonds, respectively

561 arrangement across the kernels, variation in the surface texture, relative affini-
 562 tivity of building components of almonds towards moisture and sensitivity of
 563 SWIR-HSI towards the moisture level. Similarly, increasing FFA distribu-

564 tion within the kernel can be seen across the column of figure 5b. PV was
 565 not affected until 3 months of storage, due to which first two columns of
 566 figure 5c showing low PV value. Third column of figure 5c representing the
 567 almonds stored for 4 months due to which majority of the kernels showed
 568 high PV values. From the industrial standpoint pixel wise classification may
 569 carry limited importance but it could be a good starting line for fundamental
 570 research to discover correlation between moisture distribution and initiation
 571 of hydrolytic and oxidative rancidity within the kernels.

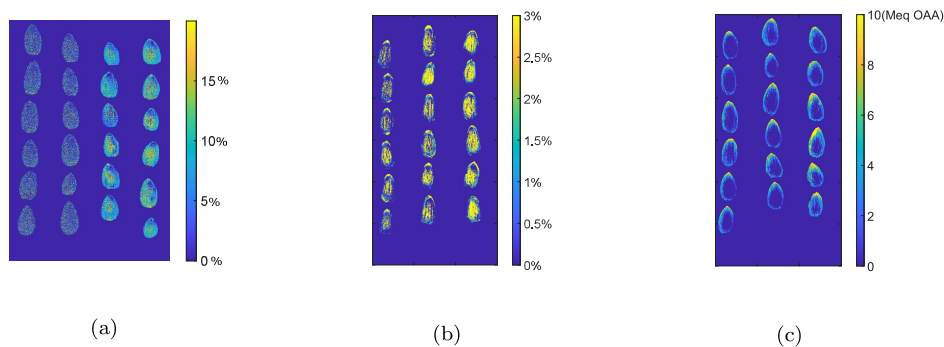


Figure 5: Visualization of response distribution for (a) Moisture (%wb), (b) FFA (%) and (c) PV (Meq OAA); units of the color bars are same as that of the corresponding responses

572 4. Conclusion

573 Current study involved combination of hyperspectral data with PLSR
 574 chemometrics technique, yielded best prediction results for moisture and
 575 rancidity of shelled almond kernels. The SG-1st derivative and SG-SNV
 576 pre-processing techniques when applied to the raw spectra showed optimal
 577 prediction results for MC and rancidity, respectively. Accuracy of the full-
 578 spectrum PLSR models were very good with R^2 and $RMSE_P$ values of 0.957,

579 0.970, 0.955 and 0.410%, 0.053%, 0.412 mEq OAA, respectively for MC,
580 FFA and PV. For rapid quantification and suitability of the system to use
581 in the commercial scale, feature wavelengths were selected using CARS-PLS
582 method without losing significant spectral information, which demonstrates
583 the novelty of the current study. Five and seven number of feature wave-
584 lengths were selected for prediction of MC and rancidity, respectively. Sim-
585 plified MLR models were developed and tested for the robustness and the
586 results showed good predictability with R^2 values of 0.941, 0.903, 0.886 and
587 $RMSEP$ values of 0.494%,0.162%, 0.658 mEq for MC, FFA and PV, respec-
588 tively. The overall results concluded that HSI system can be used for basic
589 research purpose to select the useful wavelengths or lab scale prediction of
590 the quality parameters and then multispectral imaging system with specific
591 filters can be developed to generate monochrome images for meeting the
592 need of rapid online prediction. The rapid analysis of all nuts would re-
593 duce the waste by early detection and removal of the rancid nuts from the
594 batch. However, automation of the developed technology in food industry
595 may face difficulty with the influence of physical and biological variability,
596 which affects the robustness of the features and algorithms. Thus, future
597 works will be focused on the development of diverse libraries considering cul-
598 tivar, maturity level, origin, and harvest season to bring sufficient variation
599 in the calibration dataset. Moreover, manufacturing of multispectral system
600 with filter wheels and transferring the calibration method could be done for
601 commercial use of the developed technology.

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608 **CrediT authors statement**

609 Brajesh Kumar Panda: Investigation (equal), data curation, writing origi-
610 nal draft (Lead); Gayatri Mishra: Methodology, Investigation (equal), Soft-
611 ware, validation, writing original draft (supporting); Wilmer Ariza Ramirez:
612 Data curation, Formal analysis, validation, writing-review and editing; Hye-
613 won Jung: Formal analysis, validation; Chandra B Singh: Conceptualisation,
614 supervision (equal), resources, writing-review and editing; Sang-Heon Lee:
615 Project administration, supervision (equal), writing-review and editing; Ivan
616 Lee: supervision (equal), writing-review and editing.

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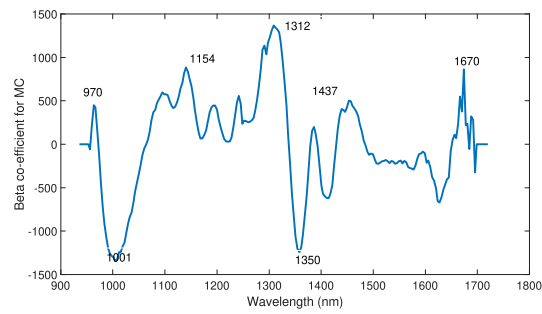
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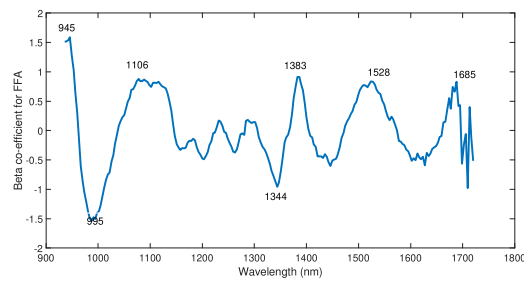
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Supplemental Materials

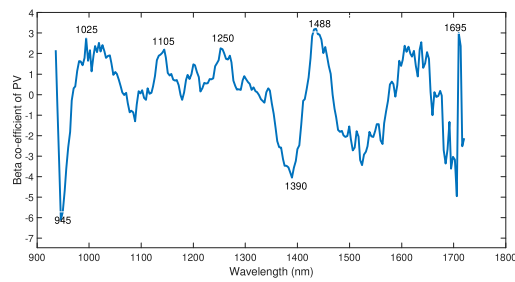
1. Beta Coefficients



(a)



(b)



(c)

Figure S1: Beta co-efficient values obtained from the PLS models developed for prediction of (a) MC (b) FFA and (c) PV of almond kernels