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TITLE PAGE
- Korean Journal for Food Science of Animal Resources -
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ARTICLE INFORMATION	Fill in information in each box below
Article Type	Research article
Article Title	Quality Assessment of Beef Using Computer Vision Technology
Running Title (within 10 words)	Assessment of Beef Quality Using Imaging Technology
Author	Md. Faizur Rahman ¹ , Abdullah Iqbal ² , Md. Abul Hashem ^{1,*} , Akinbode A. Adedeji ³
Affiliation	¹ Md. Faizur Rahman, Department of Animal Science, Bangladesh Agricultural University, Mymensingh-2202, Bangladesh; ² Dr. Abdullah Iqbal, Department of Food Technology and Rural Industries, Bangladesh Agricultural University, Mymensingh-2202, Bangladesh. ³ Dr Akinbode A. Adedeji, Department of Biosystems and Agricultural Engineering, 128 C.E. Barnhart Building, University of Kentucky, Lexington KY. 40546 USA ^{1,*} Corresponding Author: Professor Dr. Md. Abul Hashem, Department of Animal Science, Bangladesh Agricultural University, Mymensingh-2202, Bangladesh.
Special remarks – if authors have additional information to inform the editorial office	This manuscript has reviewed and edited thoroughly for technical and grammatical corrections by Dr. Akinbode A. Adedeji who is one of the Co-Author. He is an Assistant Professor of the Department of Biosystems and Agricultural Engineering, University of Kentucky, Lexington KY. 40546 USA.
ORCID (All authors must have ORCID) https://orcid.org	Md. Faizur Rahman: https://orcid.org/0000-0001-9834-5928 Abdullah Iqbal: https://orcid.org/0000-0003-3784-3522 Md. Abul Hashem: https://orcid.org/0000-0001-5691-3544 Akinbode A. Adedeji: https://orcid.org/0000-0002-3497-0166
Conflicts of interest List any present or potential conflict s of interest for all authors. (This field may be published.)	The authors declare no potential conflict of interest.
Acknowledgements State funding sources (grants, funding sources, equipment, and supplies). Include name and number of grant if available. (This field may be published.)	This research was supported by the grant provided by the Ministry of Education, The People's Republic of Bangladesh.
Author contributions (This field may be published.)	Md. Faizur Rahman: Data curation, Investigation, Writing- Original draft preparation. Abdullah Iqbal: Conceptualization, Methodology, Software Data curation. Md. Abul Hashem: Supervision, Fund Acquisition, Visualization, Investigation, Writing- Review and Editing. Akinbode A. Adedeji: Writing- Reviewing and Editing.

Ethics approval (IRB/IACUC) (This field may be published.)	This article does not contain any studies with human participants or animals performed by any of the authors. Meat samples were collected from the slaughterhouse.
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6 **CORRESPONDING AUTHOR CONTACT INFORMATION**

For the <u>corresponding</u> author (responsible for correspondence, proofreading, and reprints)	Fill in information in each box below
First name, middle initial, last name	Md. Abul Hashem
Email address – this is where your proofs will be sent	hashem_as@bau.edu.bd
Secondary Email address	hashem_mdabul@yahoo.com
Postal address	Department of Animal Science, Bangladesh Agricultural University, Mymensingh-2202, Bangladesh
Cell phone number	+880-1721310621
Office phone number	+880-091-67401-6/2633
Fax number	+88091-61510

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Abstract Imaging technique or computer vision technology has received huge attention as a rapid and non-destructive technique throughout the world for measuring quality attributes of agricultural products including meat and meat products. This study was conducted to test the ability of computer vision technology to predict the quality attributes of beef. Images were captured from *longissimus dorsi* muscle in beef at 24 hours post-mortem. Traits evaluated were color value (L^*, a^*, b^*), pH, drip loss, cooking loss, dry matter, moisture, crude protein, fat, ash, Thiobarbituric acid reactive substance (TBARS), Peroxide value (POV), Free fatty acid (FFA), Total coliform count (TCC), Total viable count (TVC) and Total yeast-mould count (TYMC). Images were analyzed using the Matlab software (R2015a). Different reference values were determined by physicochemical, proximate, biochemical and microbiological test. All determination was done in triplicate and the mean value was reported. Data analysis was carried out using the programme Statgraphics Centurion XV.I. Calibration and validation model were fitted using the software Unscrambler X version 9.7. A higher correlation found in a^* ($r = 0.65$) and moisture ($r = 0.56$) with ' a^* ' value obtained from image analysis and the highest calibration and prediction accuracy was found in Lightness ($R^2_c = 0.73$, $R^2_p = 0.69$) in beef. Results of this work show that computer vision technology may be a useful tool for predicting meat quality traits in the laboratory and meat processing industries.

Keywords beef quality, computer vision technology, correlation, calibration, validation

Introduction

Beef is a major source of essential amino acids needed in the human diet, and it attracts a premium price. Aside from being a source of protein, it is also a major source of other valuable

31 nutrients such as vitamin, fat, and micronutrients, all of which are responsible for good human
32 health. In recent years, meat quality has become a relevant topic for consumers concerning health
33 and for meat industry stakeholders because it affects their profitability (Hocquette and Chatellier,
34 2011). Meat quality is usually defined by physical, chemical and biological attributes. With the
35 current growing need for low production cost and high efficiency, the meat processing industry
36 is facing several challenges, including maintenance of high-quality standards and assurance of
37 food safety while avoiding liability issues. Meeting these challenges has become crucial in
38 regards to grading beef for different markets. Traditionally, quality assessment of beef involves
39 human visual inspection, in addition to chemical or biological determination experiments which
40 are tedious, time-consuming, destructive and sometimes environmentally unfriendly. Meat
41 processing companies and suppliers need accurate, fast, real-time, low-cost and non-chemical
42 detection technologies to optimize quality assurance of meat to enable them to satisfy different
43 market's needs, thereby raising their competitiveness and expanding their market share. Imaging
44 methods have been recently applied successfully to visually assess the quality or to classify meat
45 or food products on the processing line based on color, shape, size, surface texture features (Iqbal
46 et al., 2010; Chmiel et al., 2011; Girolami et al., 2013; Jackman et al., 2011). Computer vision
47 (CV) is one such method. It is a nondestructive, fast, cost-efficient, consistent and objective
48 method for the inspection and assessment of food quality and safety in the processing line
49 (Gümüş et al., 2011). It is an RGB color vision method that has achieved good results in the
50 assessment of the external features of foodstuffs (Tan, 2004). CV has such advantages as being
51 online, non-invasive and thus nonhazardous (Chmiel et al., 2016). As a rapid and non-destructive
52 technique, imaging technique has received huge attention in recent times for measuring quality
53 attributes of agricultural products including meat and meat products. Part of the reasons why

54 computer vision has gained popularity in times includes the fact that it can obtain reliable and
55 reproducible results (Yagiz et al., 2009). It is also able to replicable, hence, it can potentially
56 replace human vision and perception of images in meat quality assessment and safety assurance.
57 Furthermore, machine vision is capable of providing reliable descriptive data with human
58 intervention, which speed up the overall evaluation or measurement processes. Finally, it is
59 proved to be objective, effective, reliable, non-destructiverobust and capable of constant
60 recording of food samples beingexamined and the effects of processing regimes which is suitable
61 and important for further or subsequent analysis (Brosnan and Sun, 2004). The superiority of the
62 imaging technique compared to traditional analysis methods is that they allow the display and
63 overlay the distribution of the analyzed properties (Turgut et al., 2014). Computer vision system
64 has been used for colour measurement in meat by Fatih et al., (2016). Researchers used
65 computer vision technology for assessing water holding capacity in meat (Qiao et al., 2007;
66 Monroy et al., 2010; ElMasry et al., 2011). Imaging analysis of images obtained from a digital
67 camera is presently being used for assessing the external qualities of meat. However, these
68 assumptions may be possible because one previous research reported that frozen breast meat
69 with low water-holding capacity had more flat in shape during extended storage time (Lee et al.,
70 2008). Direct measurements are inconvenient and time-consuming when used in the continuous
71 processing of meat. Thus, image analysis with a digital camera may provide an alternative
72 method for evaluating or predicting the quality attributes by determining the conformation
73 parameters and surface appearance such as color, texture, bitonality etc. A number of high-
74 performance techniques have been applied successfullyfor determining or predicting the quality
75 characteristics of various meat and meat products, such as the hyperspectral imaging technique
76 (Qiao et al., 2007; Iqbal et al., 2013), near-infrared (NIR) imaging (ElMasry et al., 2011), and

77 nuclear magnetic resonance (NMR) (Bertram et al., 2001) have been used. However, these
78 techniques require costly equipment, whereas image analysis using a digital camera is
79 inexpensive. Although the machine vision has been originated during the dates back to the
80 1960s, it has not been introduced commercially in the food or processing industries until the
81 1990s. Machine vision has a distinct drawback such as its application during analysis of digital
82 images, it is restricted to the identification and extraction of external image features or quality
83 factors like color, size, and surface structure (Chmiel et al., 2011; Chmiel et al., 2012; Penman,
84 2001; Zhang et al., 2015). In consequence, it can not be used in chemometrics modeling in which
85 chemical composition and internal quality characteristics of meat or samples under consideration
86 (Peng and Dhakal, 2015).

87 Though the hypothesis of computer vision technology is related to conformation parameter of
88 foodstuffs, an effort has been taken to find out the correlation between image value and chemical
89 composition of beef through this experiment.

90 **Materials and Methods**

91 **Sample preparation**

92 Beef samples were collected from 45 carcasses between the 12th and 13th ribs (*longissimus*
93 *dorsi* muscle) of young zebu bulls. All samples were collected from different abattoir at
94 Mymensingh town, Bangladesh. The indigenous bulls were around two and a half years of age
95 with a live weight ranges from 250 to 300 kg. Each steak was 2.5 cm thick and weight was
96 around 130 gm. Samples were areal-packed and stored in the refrigerator for 24 hours at 4°C as
97 it takes at least 24 hours to convert muscle into meat. After 24 hours samples were removed from
98 the refrigerator and then kept it in a tray for about 10-12 minutes to allow moisture to appear on

99 beef surface. Then the surface of the samples were soaked gently with the help of blotting paper
100 which subsequently used for better color value estimation.

101

102 **Image acquisition**

103 Image acquisition of the sample was performed with the help of imaging system (Computer
104 Vision System, Figure 1) developed locally following the information reported by Iqbal et al,
105 (2010) and Valous et al., (2009). The main components of the developed system are: an
106 illumination source, a color digital camera (Canon IXUS, Model No. 190), and a computer-
107 supported with an image acquisition software package (Matlab 2015a, The Mathworks, Inc,
108 USA). Images of the samples were captured using the camera of imaging system and were stored
109 in the computer for further processing. An image processing software (Matlab 2015a, The
110 Mathworks, Inc, USA) was applied for image analysis.

111 **Reference analysis**

112 **1. Surface colour evaluation**

113 The surface colour of the samples were measured in terms of L* (lightness), a* (redness) and b*
114 (blueness) values using a Chroma meter (CR-400, Konica Minolta, Japan) following the
115 guidelines provided by CIE (Commission International de l'Eclairage) system (CIE,1976).

116 **2. Physico-chemical analysis**

117 **(i) pH value recording:** The pH value in beef was measured by meat pH meter (Model no.
118 HI99163, Hanna Instruments, Inc, USA). The pH meter was adjusted with pH 7.01 buffer
119 solution before the measurement. The electrodes were rinsed with cleaning solution after use.

120 **(ii) Drip Loss (DL) measurement:** For DL measurement 30 g sample was hung with a wire and
121 kept in an air tight plastic container for 24 hours. After 24 hours the sample was weighed and
122 calculated the difference. It was expressed as percentage (%).

123 **(iii) Cooking loss (CL) measurement:** 30 g beef sample was taken in a poly bag and boiled it in
124 water bath until the temperature rises to 71⁰C in sample. Beef with 71⁰C was taken out from the
125 water bath and soaked it with tissue paper. Weight loss of the sample was measured during
126 cooking beef. CL was calculated using following formula:

$$127 \text{ CL\%} = \frac{\text{Wt. before cooking} - \text{Wt. after cooking}}{\text{Wt. before cooking}} \times 100$$

128 **(iv) Proximate Analysis**

129 Moisture, protein, fat, and ash was determined as per the Standard procedures of Association of
130 Official Analytical Chemists (AOAC, 2005).

131 **3. Biochemical analysis**

132 Three types of biochemical analysis were carried out in this study: (i) Thiobarbituric Acid
133 Reactive Substance (TBARS), (ii) Free Fatty Acid (FFA) and (iii) Peroxide value
134 (POV) measurement. Three types of analysis are discussed below:

135 **(i) Thiobarbituric Acid Values (TBARS) measurement:**

136 Lipid oxidation was assessed in triplicate using the 2-thiobarbituric acid (TBA) method
137 described by Schmedes *et al* (1989). Samples (5 g) was blended with 25 mL of 20%
138 trichloroacetic acid solution (200 g/L of trichloroacetic acid in 135 mL/L phosphoric acid
139 solution) in a homogenizer (IKA) for 30 seconds . The homogenized sample was filtered with
140 Whatman filter paper number 1 and 2 mL of the filtrate was added to 2 mL of 0.02 M aqueous
141 TBA solution (3 g/L) in a test tube. The test tube was incubated at 100⁰ C for 30 minutes and
142 cooled with tap water. The absorbance was measured at 532 nm using a UV-VIS

143 spectrophotometer (UV-1200, Shimadzu, Japan). The TBA value was expressed as mg
144 malonaldehyde per kilogram of sample.

145 **(ii) Peroxide Value (POV) analysis (meq/kg):**

146 Peroxide value (POV) was determined according to Sallam *et.al.* (2004). The sample (3 g) was
147 weighed in a 250-mL glass stopper Erlenmeyer flask and heated in a water bath at 60⁰ C for 3
148 min to melt the fat, then thoroughly agitate for 3 min with 30 mL acetic acid-chloroform solution
149 (3:2 v/v) to dissolve the fat. The sample was filtered under vacuum through Whatman filter paper
150 number 1 to remove meat particles. Saturated potassium iodide solution (0.5 mL) was added to
151 filtrate and continue with addition of starch solution. The titration was allowed to run against
152 standard solution of sodium thiosulfate (25/1).

153 The formula is expressed as:

$$154 \text{ POV (meq/kg)} = \frac{S \times N}{W} \times 100$$

155 Where,

156 S is the volume of titration (mL),

157 N is the normality of sodium thiosulfate solution ($n = 0.01$) and

158 W is the sample weight (g).

159 Peroxide Value (POV) was expressed as milliequivalent peroxide per kilogram of sample.

160 **(iii) Free Fatty Acid (%) analysis:**

161 Free fatty acid value was determined according to Rukunudin *et al.* (1998). 5 g sample was
162 dissolved with 30 mL chloroform using a homogenizer (IKA T25 digital Ultra-Turrax, Germany)
163 at 10.000 rpm for 1 min. The sample was filtered under vacuum through Whatman filter paper
164 number 1 to remove meat particles. After five drops of 1% ethanolic phenolphthalein was added

165 as indicator to filtrate, the solution was titrated with 0.01N ethanolic potassium hydroxide. The
166 formula is expressed as:

$$167 \text{ FFA (\%)} = \frac{\text{Titrate required (ml)} \times \text{Normality of KOH} \times 28.2}{\text{Sample weight (g)}}$$

168 **4. Microbiological analysis**

169 Microbiological analysis was determined by Ikhlas *et al.* (2012). The procedures which were
170 followed for microbial assessment of total viable count, total coliform count and total yeast-
171 mould count, are described below:

172 **(i) Enumeration of total viable count (TVC):**

173 For the determination of total bacterial counts, 0.1 ml of each ten-fold dilution was transferred
174 and spread on triplicate PCA agar using a sterile pipette for each dilution. The diluted samples
175 were spread as quickly as possible on the surface of the plate with a sterile glass spreader. One
176 sterile spreader was used for each plate. The plates were then kept in an incubator at 35°C for 24-
177 48 hours. After incubation, 30-300 colonies were counted with the aid of a colony counter. The
178 average number of colonies in a particular dilution was multiplied by the dilution factor to obtain
179 the total viable count. The results of the total bacterial count expressed as the number of
180 organism of colony forming units per gram (CFU/g) of sample.

181 **(ii) Enumeration of total coliform count (TCC):**

182 For the determination of total coliform counts, 0.1 ml of each ten-fold dilution was transferred
183 and spread on triplicate MA agar using a sterile pipette for each dilution. The diluted samples
184 were spread as quickly as possible on the surface of the plate with a sterile glass spreader. One
185 sterile spreader was used for each plate. The plates were then kept in an incubator at 35°C for 24-
186 48 hours. After incubation, 30-300 colonies were counted with the aid of a colony counter. The
187 average number of colonies in a particular dilution was multiplied by the dilution factor to obtain

188 the total coliform count. The results of the total bacterial count expressed as the number of
189 organism of colony forming units per gram (CFU/g) of sample.

190 **(iii) Enumeration of Yeast-Mould count (TYMC):**

191 For the determination of yeast and mould counts, 0.1 ml of each ten-fold dilution was transferred
192 and spread on triplicate PDA agar using a sterile pipette for each dilution. The diluted samples
193 were spread as quickly as possible on the surface of the plate with a sterile glass spreader. One
194 sterile spreader was used for each plate. The plates were then kept in an incubator at 25°C for 48-
195 72 hours. After incubation, 30-300 colonies were counted with the aid of a colony counter. The
196 average number of colonies in a particular dilution was multiplied by the dilution factor to obtain
197 the yeast and mould count. The results of the yeast and mould count were expressed as the
198 number of organism of colony forming units per gram (CFU/g) of sample.

199 **Statistical analysis**

200 Descriptive statistical analysis and Pearson correlations between the image data and reference
201 data were both determined using the statistical package, Statgraphics Centurion XV.I.
202 STATPOINT TECHNOLOGIES, INC. Warrenton, Virginia, USA with a significance level of
203 $P < .05$. The calibration and validation model were fitted using the software Unscrambler X
204 version 9.7.

205 **Results and Discussion**

206 **Color value estimation**

207 Color measurement is more important for the visual impression of the meat than an actual
208 quality parameter. Color is usually measured in the CIE lab $L^*a^*b^*$ scale where L^* denotes the
209 brightness, a^* the redness and b^* the yellowness. The color values obtained from image analysis
210 in beef were 50.75 ± 3.43 , 13.08 ± 6.96 , 13.66 ± 2.33 for L^* , a^* , b^* respectively (Table 1). The L^* ,

211 a*, b* values from direct measurement using colorimeter were 41.67 ± 4.08 , 14.35 ± 2.1 and
212 10.44 ± 1.89 respectively and shown in Table 1. Where Fatih et al. (2016) found 48.90, 24.21 and
213 12.31 for L*,a*,b* respectively from image analysis and 46.73 ± 1.01 , 21.94 ± 1.24 and 13.11 ± 1.00
214 for L*,a*,b* respectively from direct measurement and Kamruzzaman et al.(2016) stated L*, a*,
215 b* values for beef were 47.25 ± 5.19 , 15.81 ± 2.25 and 7.56 ± 3.29 respectively. The L* and a*
216 values obtained from colorimeter by Weglarz (2010) were 37.40 ± 1.38 and 13.44 ± 2.07
217 respectively that are almost similar to the findings of the present study.

218 **Physicochemical properties**

219 The descriptive statistic of pH, drip loss and cooking loss are shown in Table 1. Measurements
220 of pH have proven to be an important analytical measurement. Muscle pH is the key to the
221 conversion of muscle to meat. During the early post-mortem changes in muscles of slaughtered
222 animals, the pH falls from around 7.0-7.2 in the muscle of a living animal to 5.5-5.8. The pH
223 value found in the LD muscle was 5.9 ± 0.12 which is close to the findings of Weglarz (2010) and
224 Rahman et al., (2015). The loss of fluid from meat is important for the industry because of its
225 economic implication. Drip loss and cooking loss obtained from the samples were 3.2 ± 0.6 and
226 29.01 ± 1.96 , respectively. The results obtained by De Marchi et al., (2007) for drip loss and
227 cooking loss as 3.87 ± 1.72 and 23.8 ± 3.55 , respectively which are very close to the present study.

228 **Proximate components**

229 Dry matter, moisture, crude protein, fat and ash content of the sample has been shown in Table
230 1. The values were 24.79 ± 2.26 , 75.21 ± 2.27 , 20.88 ± 2.69 , 1.57 ± 0.63 , 1.28 ± 0.21 for dry matter,
231 moisture, crude protein, fat, and ash, respectively. These findings are in close agreement with
232 those reported by De Marchi et al., (2007).

233

234 **Biochemical properties**

235 The oxidative stability of beef depends upon the balance of anti and pro-oxidants and the
236 composition of these oxidation substrates (Bertelsen et al., 2000). Average TBARS, peroxide and
237 free fatty acid value of beef *longissimus dorsi* were found 0.11 ± 0.01 , 1.85 ± 0.35 and 0.04 ± 0.01
238 respectively (Table 1) where researchers found almost same values in fresh beef from hind limb
239 of bull (Rahman et al., 2000). The biochemical traits measured in this experiment were lower
240 than the limit (TBARS: <0.6 mgMDA/kg, POV: <6 meq/kg and FFA <1.2) for rancidity.

241 **Microbiological analysis**

242 Microbiological traits measured by the laboratory method are presented in Table 1. The average
243 values with standard deviations were 5.09 ± 0.05 , 5.91 ± 0.07 and 7.7 ± 0.08 for TCC, TYMC, and
244 TVC respectively. The TVC level of this study was in close agreement with the findings of Saba
245 et al., (2018) but the TCC and TYMC value were higher than results of Murshed et al., (2016),
246 Afrin et al., (2017) and Alam et al., (2017). The possible cause of this variation in microbial load
247 might be due to the differences in the aging period.

248 **Correlation between computer vision technology and conventional analytical technology**

249 Correlation between image data and reference data of beef *longissimus dorsi* is presented in
250 Table 2. The L^* value from image analysis had medium correlation with L^* (0.46), pH (0.41),
251 DL% (0.37), DM (0.31), CP (0.24) and Ash (0.25) obtained from laboratory method. A higher
252 correlation found in a^* (0.65) and moisture (0.56) with ' a^* ' value obtained from imaging
253 technique. The ' b ' value resulted from image analysis had a medium correlation with fat (0.33)
254 and TVC (0.37) whereas Mello et al., (2015) measured intramuscular fat and Luo et al., (2018)
255 determined marbling in beef using image processing. The imaging technique, computer vision
256 system was very similar that were used for both the previous works and present study.

257 **Prediction of beef longissimus dorsi quality traits**

258 It is known that the coefficient of determination (R^2) indicates the accuracy of model, varying
259 from 0 to 1. Table 3 presents the results of calibration and prediction of color, pH, drip loss, dry
260 matter, crude protein, peroxide value and ash content of the meat samples using image data. The
261 calibration coefficients, R^2 range from 0.21-0.73 and the range of prediction data R^2 is 0.01-0.69.
262 The root mean square error for calibration and prediction are general low, less than 2.4.
263 Lightness, L^* has the most association with the image data in both calibration and prediction
264 analyses. ElMasry et al., (2012) predicted L^* and pH values with coefficients of determination
265 (R^2) of 0.88 and 0.73 and root mean square errors estimated by cross validation (RMSECV) of
266 1.21 and 0.06 respectively in beef. Chmiel et al., (2011) used computer image analysis to detect
267 PSE defects in pork meat. They found higher values of L^* (56.01 ± 1.62) in PSE meat compared
268 with normal meat (48.44 ± 0.52). From image analysis they found significantly higher values of R,
269 G, B components for PSE meat compared with normal meat. Sun et al. (2016) applied computer
270 vision method for assessing the color score in pork meat. They found very significant correlation
271 ($p < 0.0001$) between L^* , a^* and b^* features of images and Minolta colorimeter 0.91, 0.80 and 0.66
272 respectively. They assessed the coefficient of determination (R^2) for predicted pork color features
273 was 0.83. The coefficient of determination was found 0.85 for CD large in IMF feature of
274 longissimus dorsi muscle in beef using an image processing algorithm by Du et al. (2008).

275 **Conclusion**

276 The study aimed at assessing the ability of the computer vision system to predict beef quality
277 traits. Samples were analyzed for color, physicochemical, proximate, biochemical and
278 microbiological values by conventional analytical techniques. Computer vision technology has
279 been standardized. After the standardization correlation coefficient was determined between

280 image data and reference data. The calibration model and validation model was applied to find
281 the level of accuracy of the computer vision technology. The highest calibration and prediction
282 accuracy was found for colour Lightness (L^*), medium accuracy was found in the redness (a^*),
283 pH, drip loss, crude protein and ash content of the sample. However, more samples and trials are
284 to be conducted in future for the development of robust models and getting higher predicting
285 values of the studied parameters.

287 **Conflicts of Interest**

288 The authors declare no potential conflict of interest.

289 **Acknowledgements**

290 This research was supported by the grant provided by the Ministry of Education, The People's
291 Republic of Bangladesh.

292 **Author Contributions**

293 Md. Faizur Rahman: Data curation, Investigation, Writing- Original draft preparation. Abdullah
294 Iqbal: Conceptualization, Methodology, Software Data curation. Md. Abul Hashem: Supervision,
295 Fund Acquisition, Visualization, Investigation, Writing- Review and Editing. Akinbode A.
296 Adedeji: Writing- Reviewing and Editing.

297 **Ethic Approval**

298 This article does not contain any studies with human participants or animals performed by any of
299 the authors. Meat samples were collected from the slaughterhouse.

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411 **Table 1**

412 Range, mean, standard deviation (SD) and coefficient of variation (CV) of beef *Longissimus*
 413 *dorsi* muscle quality traits

Attribute	n	Range	Mean	SD	CV%
L* _{image}	45	38.37-56.18	50.75	3.43	6.76
a* _{image}	45	1.57-23.26	13.08	6.96	53.19
b* _{image}	45	6.71-18.09	13.66	2.33	17.08
L*	45	37.18-49.72	41.67	4.08	9.79
a*	45	10.48-17.71	14.35	2.1	14.65
b*	45	8.14-15.24	10.44	1.89	18.08
pH	45	5.8-6.1	5.9	0.12	2.08
DL%	45	2.37-4.73	3.2	0.6	18.6
CL%	45	24.8-31.9	29.01	1.96	6.75
DM%	45	21.22-28.98	24.79	2.26	9.12
Moisture%	45	71.02-78.98	75.21	2.27	3.02
CP%	45	17.5-26.35	20.88	2.69	12.87
EE%	45	0.72-3.5	1.57	0.63	40.05
Ash%	45	0.88-1.67	1.28	0.21	16.1
TBARS	45	0.09-0.13	0.11	0.01	9.21
POV	45	0.99-2.65	1.85	0.35	18.8
FFA	45	0.02-0.06	0.04	0.01	26.37
TCC	45	4.93-5.16	5.09	0.05	0.98
TYMC	45	5.71-6.01	5.91	0.07	1.26
TVC	45	7.48-7.87	7.7	0.08	1.09

414 L*_{image}, L*value from imaging analysis; a*_{image}, a* value from image analysis; b*_{image}, b*
 415 value from imaging analysis; n, sample size; SD, standard deviation; CV, co-efficient of
 416 variation; DL, drip loss; CL, cooking loss; DM, dry matter; CP, crude protein; EE, ether extract;

417 TBARS, thiobarbituric acid reactive substance; POV, per-oxide value; FFA, free fatty acid; TCC,
418 total coliform count; TYMC, total east-mould count; TVC, total viable count.

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420 **Table 2**

421 Pearson correlation coefficients between image data and reference data for quality attributes
 422 of beef *longissimusdorsi* muscle

	L*_{image}	a*_{image}	b*_{image}
L*	0.46 ^{***}	-0.86 ^{***}	0.05
a*	-0.24	0.65 ^{***}	0.1
b*	0.05	0.29	-0.1
pH	0.41 ^{**}	-0.75 ^{***}	0.21
DL%	0.37 ^{**}	-0.78 ^{***}	0.29
CL%	-0.1	0.26	-0.04
DM	0.31 [*]	-0.56 ^{***}	0.28
Moisture	-0.31 [*]	0.56 ^{***}	-0.28
CP	0.24	-0.69 ^{***}	0.16
EE	0.08	-0.17	0.33 [*]
Ash	0.25	-0.72 ^{***}	-0.03
TBARS	0.01	-0.21	-0.15
POV	0.1	-0.43 ^{**}	0.03
FFA	-0.06	0.11	-0.21
TCC	-0.1	0.14	-0.05
TYMC	-0.16	0.23	0.16
TVC	-0.05	-0.1	0.37 ^{**}

423 L*_{image}, L* value from image analysis; a*_{image}, a* value from image analysis; b*_{image}, b*
 424 value from imaging analysis; DL, drip loss; CL, cooking loss; DM, dry matter; CP, crude protein;
 425 EE, ether extract; TBARS, thiobarbituric acid reactive substance; POV, per-oxide value; FFA,
 426 free fatty acid; TCC, total coliform count; TYMC, total east-mould count; TVC, total viable
 427 count *p<0.05, **p<0.01, ***p<0.001.

428

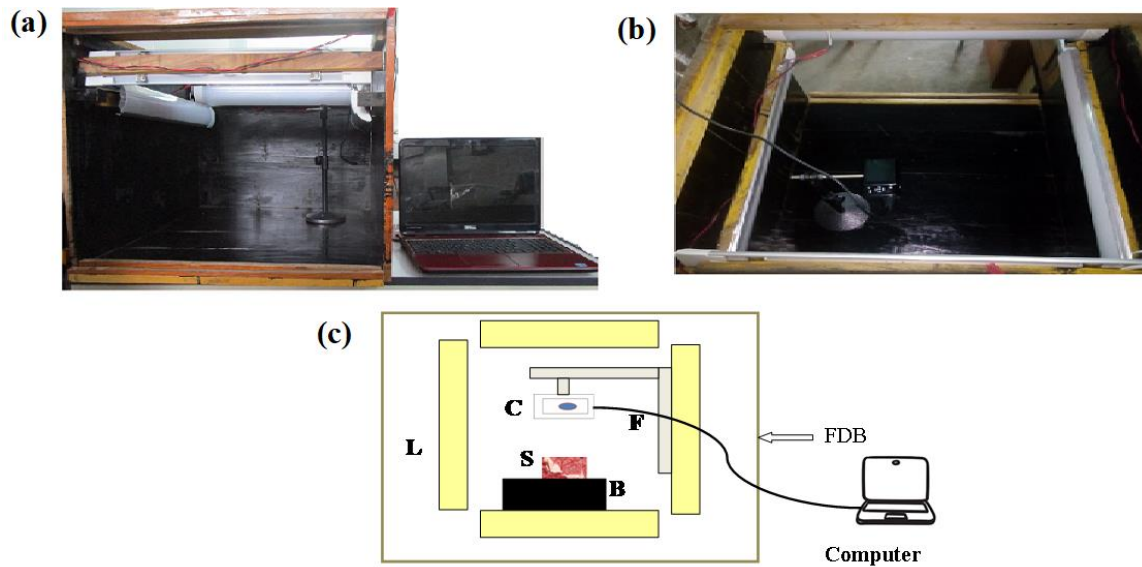
429 **Table 3**430 Best fitting predictions of ten quality traits on beef *longissimus dorsi* using image technology

Variable	n	R²_C	RMSE_C	R²_P	RMSE_P
L*	45	0.73	2.09	0.69	2.23
a*	45	0.47	1.52	0.34	1.69
b*	40	0.21	2.02	0.01	2.36
pH	45	0.59	0.07	0.52	0.08
DL	45	0.67	0.34	0.6	0.37
DM	45	0.37	1.77	0.26	1.93
Moisture	45	0.37	1.77	0.26	1.93
CP	45	0.53	1.83	0.43	2.00
Ash	45	0.56	0.14	0.45	0.15
POV	45	0.22	0.3	0.06	0.33

431 DL, drip loss; DM, dry matter; CP, crude protein; POV, per-oxide value; n, number of samples;
 432 R²_C, coefficient of determination of calibration; RMSE_C, root mean square error of calibration;
 433 R²_P, coefficient of determination of prediction; RMSE_P, root mean square error of prediction.

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437 **Figure 1. Computer vision system developed in the Laboratory: (a) Front view; (b) Top**
438 **view and (c) Schematic diagram with its components: L= Light sources, C=Camera,**
439 **S=Sample, B= Black background, F= Attachment for Camera, FDB= Frame with dark box.**

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