

1                   **RUNNING HEAD: Breed effect on nutritional value of beef**

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3           **Fatty acid profile, mineral content and palatability of beef from a multibreed Angus-**

4                                   **Brahman population**

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15 **ABSTRACT:**

16 Consumers demand meat that is both healthy and palatable, two attributes of meat that  
17 are affected by lipid content. Many cattle in the southern US are *Bos indicus* influenced, as this  
18 improves the ability to survive and thrive in these subtropical regions. However, these animals  
19 tend to have leaner carcasses and less marbled meat products. Thus, the objective of this study  
20 was to examine the effect of percent Brahman genetics on carcass characteristics, palatability,  
21 fatty acids profile and minerals content in LM of steers from a multibreed population ranging  
22 from 100% Angus to 100% Brahman. Breed effect was significant for birth weight ( $P = 0.0003$ ),  
23 weaning weight ( $P < 0.0001$ ), HCW ( $P < 0.0001$ ), dressing percentage ( $P = 0.0008$ ), ribeye area  
24 ( $P = 0.002$ ), quality grade ( $P < 0.0001$ ), and marbling score ( $P < 0.0001$ ), and all these traits  
25 except dressing percentage decreased as the percentage of Brahman increased. Among  
26 palatability traits, breed group had a significant effect only on tenderness (TEND) and  
27 connective tissue (CT) scores ( $P < 0.0001$ ). Least squares means decreased from Angus ( $5.75 \pm$   
28  $0.13$  TEND score and  $6.29 \pm 0.14$  CT score, respectively) to Brahman ( $4.84 \pm 0.10$  TEND score  
29 and  $5.49 \pm 0.11$  CT score, respectively) as indicated by a significant linear effect. Breed group  
30 significantly affected the percentage of several individual fatty acids, saturated fatty acids (SFA)  
31 and polyunsaturated fatty acids (PUFA), but not monounsaturated fatty acids (MUFA). The  
32 100% Angus group had the highest percentage of SFA at 49.92%, which was significantly higher  
33 ( $P < 0.05$ ) than the SFA percentage in the 50%, 75%, and 100% Brahman breed groups. Brangus  
34 animals also had an increased SFA percentage compared to the 100% Brahman animals ( $P <$   
35  $0.05$ ). No significant effect was identified for the concentration of PUFA across the six breed  
36 groups ( $P = 0.14$ ). Least squares means decreased from 100% Angus to 100% Brahman for  
37 concentration of total fat, SFA and MUFA (g/mg meat). The concentration of magnesium ( $P <$

38 0.0001), phosphorus ( $P = 0.06$ ) and potassium ( $P = 0.06$ ) increased as the percentage of  
39 Brahman increased. Our study shows that breed has a significant effect on the fatty acid profile  
40 of beef. Cattle with high Brahman percentage, which are characterized by lower marbled meat,  
41 will present a more favorable healthfulness profile with reduced content of SFA and MUFA but  
42 the same content of PUFA as purebred Angus animals.

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44 **Key words:** beef cattle, *Bos indicus*, healthfulness, mineral content

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## INTRODUCTION

47 Brahman genetics are extensively used in crossbreeding programs in the southeastern  
48 regions of the United States (Cundiff et al., 2012; Lamy et al., 2012), characterized by hot and  
49 humid conditions typical of tropical and subtropical environments. While Brahman cattle are  
50 well known for their adaptability in subtropical climates (Hansen, 2004), they tend to produce  
51 less marbling within the final beef product (Johnson et al., 1990; Pringle et al., 1997; Elzo et al.,  
52 2012). Several studies addressing relevant economic traits such as growth, carcass and  
53 reproduction in *Bos indicus* influenced cattle prevalent in the southern US have been published  
54 (Riley, 2002; Riley et al., 2012; Elzo et al., 2014; Elzo et al., 2016; Elzo et al., 2017; Leal-  
55 Gutiérrez et al., 2018), but information regarding fatty acid composition and mineral content are  
56 scarce.

57 A recent Beef Demand study (2013, Schroder et al.) identified seven factors as  
58 consequential for driving beef demand. Ranked in the order of their relevance to consumers these  
59 factors are beef price, food safety, product quality, health, nutrition, social aspects, and  
60 sustainability. Given that the industry cannot control price, the report identified food safety,

61 product quality, nutritional value and healthfulness as the key attributes that the industry can and  
62 should focus on. Beef consumption helps Americans fulfill their daily-recommended dietary  
63 intake of protein, by providing 20g of protein per 100g of beef consumed, while also providing  
64 many nutrients with positive effects on human health (mono (MUFA) and polyunsaturated fatty  
65 acids (PUFA), such as omega-3 and conjugated linoleic acid (CLA), iron, zinc, vitamin B6, etc.).  
66 However, beef is also associated with characteristics that are often perceived as negative, such as  
67 high levels of saturated fatty acids (SFA) and high caloric content. Knowledge of the role of *Bos*  
68 *indicus* influenced cattle concerning these areas is essential for the prosperity of the beef industry  
69 in the southern US.

70 Intramuscular fat depot or “marbling” is a key factor in determining carcass value. While  
71 *Bos taurus* breeds of cattle, such as Angus, are known for their superior marbling potential, *Bos*  
72 *indicus* breeds have the tendency to produce less marbled beef products. The amount of  
73 intramuscular fat described by marbling and the fatty acid composition determines the  
74 healthfulness value of the beef product. Polyunsaturated (PUFA) and monounsaturated fatty  
75 acids (MUFA) are known to have cholesterol-lowering properties, and reduce the risk of  
76 coronary vascular disease among other healthful attributes. On the contrary, several short chain  
77 saturated fatty acids (SFA) are associated with increased risk of coronary vascular disease  
78 (Bonanome and Grundy, 1988; Derr et al., 1993; Judd et al., 2002; Brouwer et al., 2010).

79 Breed, along with other factors such as sex, age and diet (Wood et al., 2008; Mateescu,  
80 2015), has an impact on both the amount of marbling and the fatty acid composition. Numerous  
81 recent reports are available regarding the fatty acid composition of *Bos taurus* breeds (Nogi et  
82 al., 2011; Xu et al., 2013; Buchanan et al., 2015; Ekine-Dzivenu et al., 2017; Zhu et al., 2017),  
83 but information on the quality of fat and the healthfulness and nutritional value of beef from *Bos*

84 *indicus* influenced cattle, in particular Brahman, is scarce (Dinh et al., 2010; Campbell et al.,  
85 2016).

86 The objective of this study was to characterize the carcass and palatability traits, fatty  
87 acid composition, and mineral content in a multibreed cattle population typical to the southern  
88 US and estimate the effect of breed composition on nutritional and healthfulness value of beef.

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## MATERIALS AND METHODS

### *Animals and Management*

92 The research protocol was approved by the University of Florida Institutional Animal  
93 Care and Use Committee number 201003744. Cattle used in this study were from the University  
94 of Florida multibreed Angus-Brahman herd (Elzo et al., 2014). A total of 230 steers across six  
95 breed groups based on the percentages of Angus breed composition were used from this herd:  
96 Angus = 100 to 80% (n = 39); 75% Angus = 79 to 65% (n = 33); Brangus = 62.5% (n = 30); 50%  
97 Angus = 59 to 40% (n = 42); 25% Angus = 39 to 20% (n = 27); Brahman = 19 to 0% (n = 59).  
98 Steers born in 2014 and 2015 were transported to a contract feeder (Quincey Farms, Chiefland,  
99 FL) where they were provided a standard feedlot diet consisting of corn, protein, vitamins, and  
100 minerals until they reached a subcutaneous fat thickness over the ribeye of approximately 1.27  
101 cm assessed thorough ultrasound. The concentrate diet had, on the average, 89.7% of DM, 14.4%  
102 of CP, 1.5 Mcal/kg DM of NEm, and 1.1 Mcal/kg DM of NEg. As cattle achieved appropriate  
103 degree of back fat thickness, they were transported to a commercial packing plant where they  
104 were harvested under USDA FSIS inspection. Steers were harvested in groups of 15-25 animals  
105 and the average slaughter age was 18.76±1.13 mo.

### *Carcass Evaluation and Sample Collection*

107 At 24 hours postmortem, carcasses were ribbed between the 12<sup>th</sup> and 13<sup>th</sup> rib, per industry  
108 standard and carcass measurements were evaluated for each animal: HCW, dressing percentage  
109 (**DP**), marbling score (**MS**; 100 to 199 = practically devoid, 200 to 299 = traces, 300 to  
110 399=slight, 400 to 499=small, 500 to 599=modest, 600 to 699=moderate, 700 to 799=slightly  
111 abundant, 800 to 899=moderately abundant, and 900 to 999=abundant), ribeye area (**REA**), and  
112 fat over the eye (**FOE**). **USDA quality grades** and yield grades were calculated according to  
113 industry standards. **Following carcass evaluation, two 2.54cm thick steaks were removed from**  
114 **the anterior end of the carcass ribbing and transferred to the University of Florida Meat**  
115 **Processing Center (Gainesville, FL). Steaks were wet aged for 14 days, then frozen (-20 °C) until**  
116 **subsequent fatty acid, mineral, tenderness, and sensory analysis.**

#### 117 *Warner-Bratzler Shear Force and Sensory Panel Analysis*

118 **One of the two frozen steaks from each animal was thawed at 3°C for 24 hours and**  
119 **cooked on an open-top, electric grill. Steaks were cooked to an internal temperature of 71°C,**  
120 **equivalent to a medium degree of doneness. Internal temperature was monitored using copper-**  
121 **constant thermocouples (Omega Engineering Inc., Stamford, CT) located in the geometric center**  
122 **of each steak. Temperature was recorded by a 1100 Labtech Notebook Pro Software version 12.1**  
123 **(Computer Boards Inc., Middleboro, MA). Once cooked, steaks were chilled at 3°C for 24 hours.**  
124 **After chilling, six 1.27 cm cores were removed from each steak parallel to the muscle fibers.**  
125 **Each of the six cores were sheared through the center (crosshead speed of 200 mm/min) with a**  
126 **Warner-Bratzler shear force (**WBSF**) head attached to a 490 N load cell using an Instron**  
127 **Universal Testing Machine (Instron Corporation, Canton, Massachusetts, USA).**

128 **The other frozen steak from each animal was handled and cooked in the same manner as**  
129 **the WBSF samples. Once cooked, steaks were cut into 1.27 cm cubes and served warm to trained**

130 sensory panelists. The sensory panel consisted of 7 to 11 trained members (AMSA, 1995) who  
131 evaluated each sample for various meat palatability traits. Sensory panel measurements analyzed  
132 by the sensory panelists included: tenderness score (**TEND**; 1=extremely tough to 8=extremely  
133 tender), connective tissue score (**CT**; 1=abundant amount to 8=none detected), juiciness score  
134 (**JUIC**; 1=extremely dry to 8=extremely juicy), beef flavor score (**FLAV**; 1=extremely bland to  
135 8=extremely intense), and off-flavor score (**OFLAV**; 1=extreme off- flavor to 6=none detected).

### 136 *Fatty Acid Extraction and Gas Chromatography Analysis*

137       After trimming external fat and connective tissue, a thin shaving across the entire steak  
138 surface was removed from each steak sample and powdered in liquid nitrogen to obtain a  
139 homogenized sample of the steak. Fatty acid extraction and analysis was performed at the W. M.  
140 Keck Metabolomics Research Laboratory, Iowa State University (Ames, IA). About 200 mg of  
141 finely ground steak samples was extracted into 1mL of 2:1 Chloroform: Methanol mixture. The  
142 extracted fats were trans-esterified with 25% Sodium Methoxide in methanol. The resulting Fatty  
143 Acid Methyl Esters (FAMES) were extracted into hexane and detected on Agilent 7890A GC-  
144 FID instrument. One microliter of the sample was injected into an Agilent 7890. A Gas  
145 Chromatograph equipped with a flame ionization detector was used for separation and  
146 quantification of the FAMES. The analysis was performed on Agilent CP-Wax 52CB column  
147 (15m, 0.32mm, 0.5um). The oven temperature program was as follows. Initial temperature of  
148 100°C, increased to 170°C with a ramp of 2°C/min, increased to 180°C with a ramp of  
149 0.5°C/min, to a final temperature of 250°C with a ramp of 1°C/min and held for 3min. The inlet  
150 temperature was 250°C and detector temperatures were 220°C. Helium was used as the carrier  
151 gas. Supelco 37 FAME mix (Catalog # CRM47885 SUPELCO) was used to generate the  
152 calibration curve for identification and quantification of FAMES.

153 Twenty eight individual fatty acids and three groups of fatty acids based on saturation  
154 level (SFA, MUFA, PUFA) were calculated and expressed as percentage of the total fatty acids  
155 and as mg/g of tissue.

#### 156 *Mineral concentrations*

157 Mineral content of LM samples were determined by inductively coupled plasma-optical  
158 emission spectroscopy (ICP-OES, SPECTRO Analytical Instruments, Mahwah, NJ). The  
159 samples were dried at 105°C for 18 to 20 h according to AOAC official method 934.01 (Davis  
160 and Lin, 2005), and moisture content was calculated. Dried samples were subjected to a closed-  
161 vessel microwave digestion process (CEM, MDS-2000, Matthews, NC) with 5 mL concentrated  
162 nitric acid and 2 mL 30% hydrogen peroxide according to AOAC official methods 999.10  
163 (Jorhem and Engman, 2000). The microwave was programmed as follows: 250 watts for 5 min,  
164 630 watts for 5 min, 500 watts for 20 min, and 0 watts for 15 min. Digested samples were  
165 transferred to 25 mL volumetric flasks and diluted with deionized water. The concentrations of  
166 iron, phosphorus, potassium, sodium, magnesium and zinc were then measured by ICP-OES.

#### 167 *Statistical Analysis*

168 All statistical analyses were performed using SAS 9.4 (SAS Inst. Inc., Cary, NC). The  
169 MEANS procedure was used to produce descriptive statistics for fatty acid composition data.  
170 Traits were analyzed using the MIXED procedure of SAS. Models for all traits included year of  
171 birth as a random effect and breed as fixed effects. Breed-group least squares means were  
172 separated using LSMEANS with the PDIFF option. To estimate the linear and quadratic effect of  
173 percent Brahman genetics, the breed groups were recoded as 0, 1, 1.5, 2, 3 and 4 indicating 0%,  
174 25%, 37.25%, 50%, 75%, and 100% Brahman genetics. The model included year of birth as a  
175 random effect and the linear and quadratic breed as a covariate. When the quadratic effect was



176 not significant, it was dropped and a model including just the linear effect was used. The  
177 intercept from this model estimates the effect of 100% Angus genes (adjusted for random year of  
178 birth effect) and the estimated regression coefficients represent the effect of replacing 25% of  
179 Angus genes by Brahman genes.

180

## 181 RESULTS AND DISCUSSION

182 Table 1 presents summary statistics for traits evaluated in this study in a population of  
183 230 animals with breed composition ranging from 100% Angus to 100% Brahman.

### 184 *Carcass Characteristics*

185 Least squares means of carcass measurements for the six breed groups in this study are  
186 presented in Table 2. The carcass data for the total 230 animals is representative of industry  
187 average quality and yield grades (Shackelford et al., 2012) and similar to previously reported  
188 data on this multibreed population (Elzo et al., 2012; Elzo et al., 2014; Elzo et al., 2016). Breed  
189 effect was significant for BWT ( $P = 0.0003$ ), WWT ( $P < 0.0001$ ), HCW ( $P < 0.0001$ ), DP ( $P =$   
190  $0.0008$ ), REA ( $P = 0.002$ ), QG ( $P < 0.0001$ ), and MS ( $P < 0.0001$ ). No breed differences were  
191 identified for FOE which was expected as a direct consequence of animals slaughtered at a  
192 similar fat thickness end point. Cattle with the highest percent Brahman (Brahman breed group)  
193 had the lowest HCW and REA compared to all other breed groups. This is in agreement with  
194 previous studies which reported Angus cattle had heavier HCW and greater REA than Brahman  
195 (Peacock et al., 1979; Lunt et al., 1985; Williams et al., 2010, Elzo et al., 2012).

196 Marbling score decreased from Angus ( $464.12 \pm 13.12$  units) to Brahman ( $352.85 \pm$   
197  $10.34$  units), with the highest percentage Brahman animals having statistically lower marbling  
198 scores than all the other breed groups ( $P < 0.0001$ ). This was also reflected in the QG which

199 followed the same trend across breed groups with Brahman having the lowest QG ( $547.37 \pm 5.6$ )  
200 compared to all other breed compositions. It is important to point out that the Angus, 75% Angus  
201 and Brangus breed groups had a marbling score higher than the industry average ( $449 \pm 94.8$ ),  
202 and the Brahman breed group was within one standard deviation of this average (Shackelford et  
203 al., 2012).

204 **Table 2** also contains least squares means for the six breed groups for meat palatability  
205 traits. **No significant differences were found for beef flavor or off flavor across the breed groups.**  
206 Breed group had a significant effect only on TEND and CT traits recorded during the trained  
207 sensory panel ( $P < 0.0001$ ). **A negative linear effect was significant with least squares means**  
208 **decreasing from Angus** ( $5.75 \pm 0.13$  TEND score and  $6.29 \pm 0.14$  CT score, respectively) to  
209 Brahman ( $4.84 \pm 0.10$  TEND score and  $5.49 \pm 0.11$  CT score, respectively). Previous reports  
210 consistently identified animals with high Brahman influence to have lower sensory tenderness  
211 scores and higher connective tissue scores (Johnson et al., 1990; Pringle et al., 1997; Elzo et al.,  
212 2012). **The lower tenderness of Brahman cattle has been attributed to increased postmortem**  
213 **calpastatin activity (Wheeler et al., 1990; Shackelford et al., 1991; Pringle et al., 1997) which**  
214 **results in a reduction in desmin and troponin-T degradation (Phelps et al., 2017).** **There are two**  
215 **important points related to the tenderness qualities of steaks from Angus, Brahman and their**  
216 **crossbreeds evaluated by WBSF or through the sensory panel. When using the WBSF, considered**  
217 **an objective measure of tenderness, no significant differences were detected along the breed**  
218 **composition continuum from purebred Angus to purebred Brahman. The coefficient of variation**  
219 **obtained from a regression analysis of sensory tenderness using WBSF values was 13.34**  
220 **indicating variation in the sensory tenderness which is not captured by the WBSF. However, a**  
221 **statistically significant difference was found when tenderness was evaluated either by the TEND**

222 or CT score during the trained sensory panel. Among other factors, tenderness as perceived by  
223 consumers is determined by the amount and solubility of the connective tissue and the amount of  
224 marbling. Marbling has a major effect on the perceived juiciness and a high correlation was  
225 reported between tenderness and juiciness (Mateescu et al., 2015), where steaks with high  
226 marbling score have superior juiciness and overall liking (Killinger et al., 2004; Okumura et al.,  
227 2007; Legako et al., 2015). It is however important to highlight the fact that although statistically  
228 significant, these differences are insignificant in terms of changes in tenderness that can be  
229 perceived by regular, untrained consumers (Watson et al., 2008). These results indicate that meat  
230 from *Bos indicus* influenced cattle tends to be less tender compared to meat from Angus animals,  
231 but these differences, when measured objectively using WBSF, are not statistically significant.  
232 Moreover, when considering that the minimum difference the average consumer can detect is a  
233 0.5 kg difference in WBSF when consuming meat at home (Miller et al., 1995), the differences  
234 found in this study are not sufficiently large to be detectable by consumers. These results suggest  
235 that selection programs aimed at improving tenderness in these *Bos indicus* populations are  
236 producing the intended results and the historical perception of tougher beef associated with  
237 *indicine* influenced cattle needs to be reevaluated.

### 238 ***Breed Effect on Fatty Acid Composition***

239 Percentages of the three main fatty acids categories in the overall population were as  
240 expected, with approximately 47.52 to 49.92% of the total lipid content being represented by  
241 SFA, 45.10 to 46.10% by MUFA, and 3.98 to 7.38% PUFA (Pitchford et al., 2002; Daley et al.,  
242 2010; Garmyn et al., 2011). The fatty acid composition in steers from six breed groups,  
243 expressed both as percentage of total fatty acids and as mg/100 g muscle, is presented in Tables 3  
244 and 4, respectively.

245 Breed group significantly affected the percentage of several individual fatty acids, SFA,  
246 and PUFA, but not MUFA. The Angus group had the highest percentage of SFA at 49.92%,  
247 which was significantly higher than the SFA percentage in the 50%, 75% and 100% Brahman  
248 breed groups ( $P = 0.006, 0.008$  and  $0.002$ , respectively). Brangus animals also had a significantly  
249 ( $P = 0.04$ ) higher SFA percentage compared to the 100% Brahman animals. These differences  
250 were mainly a consequence of higher percentages of C16:0 and C18:0 in Angus or Brangus  
251 compared to animals with higher Brahman composition.

252 The opposite relationship was observed for PUFA, with the percentage of these class of  
253 fatty acids increasing as the percentage of Brahman genetics increased. Meat from Brahman  
254 steers had the highest percentage of PUFA ( $7.38 \pm 0.39$ ) compared to all other breed groups,  
255 while steers from the Angus group had a significantly lower percentage of PUFA compared to  
256 the 50%, 75% and 100% Brahman breed groups.

257 The fatty acid concentration (mg/g meat) showed a significant breed group effect for  
258 most of the individual fatty acids and for total fat, SFA, and MUFA. No significant effect was  
259 identified for the concentration of PUFA across the six breed groups ( $P = 0.14$ ). Least squares  
260 means showed a downward trend from Angus to Brahman for concentration of total fat, SFA and  
261 MUFA. For all three fatty acid categories, Angus concentrations were significantly higher than  
262 the concentrations in the 50%, 75%, and 100% Brahman breed groups, and the concentration in  
263 the 100% Brahman steers was significantly lower than steers from any other breed group.

264 In this study, the LM from Angus cattle was found to have 50% greater amount of  
265 intramuscular fat than Brahman cattle. The lower fat content of animals from the Brahman breed  
266 group is in agreement with many previous reports (Wood et al., 2008; Mateescu, 2015)  
267 describing a lower intramuscular fat content of *Bos indicus* animals. This greater accumulation of

268 fat in Angus animals was achieved by approximately 50% increase in total SFA and MUFA  
269 concentration compared to Brahman. This is to be expected, as triacylglycerols deposited into  
270 adipocytes are mostly SFA and MUFA from dietary sources and de novo synthesis (Jenkins,  
271 1994; Scollan et al., 2014), while the concentration of phospholipids which are rich in PUFA  
272 remains constant and relatively independent from the total fat amount. However, several  
273 important trends with respect to the healthfulness of the beef products from different breed  
274 groups are emerging. On a percentage basis, steaks from animals with a high Brahman  
275 percentage had significantly lower SFA and significantly higher PUFA, which suggests a higher  
276 healthfulness value of steaks from Brahman cattle. More importantly, the lower SFA is due  
277 mostly to a decrease in the percent of short chain SFA (C10:0, C11:0, C12:0 and C16:0) while  
278 C18:0 showed no significant difference among breed groups. Palmitic acid (C16:0) is a saturated  
279 fatty acid accounting for about 27% of the fatty acids in beef and has been shown to raise serum  
280 cholesterol levels (Grundy, 1994) predominantly by increasing the LDL cholesterol levels. This  
281 fatty acid accounts for most of the cholesterol-raising activity from beef, thereby increasing the  
282 risk of atherosclerosis, cardiovascular disease, and stroke (Brouwer et al., 2010). On the other  
283 hand, stearic acid (C18:0) accounts for about 18% of the fatty acid in beef. Its effect on total  
284 cholesterol is minimal and not detrimental to human health (Bonanome and Grundy, 1988; Zock  
285 and Katan, 1992; Derr et al., 1993; Judd et al., 2002). For practical purposes, stearic acid is  
286 essentially neutral in its effects on serum total cholesterol, similar to C18:1 or oleic acid  
287 (Grundy, 1994).

288 The second and more important finding is related to the fatty acid concentration on a  
289 mg/g meat basis. Most studies on fatty acid composition in beef cattle have reported the  
290 normalized percentage of total fatty acids, which describes the lipid quality and is driven by

291 strong relationships among fatty acids. Steers with a high Brahman breed composition have  
292 leaner meat compared to purebred or high Angus percentage steers and this is accompanied by a  
293 high ratio of phospholipid to triacylglycerol in the fat fraction (Scollan et al., 2007; Buchanan et  
294 al., 2015). This is expected and it is reflected in the significantly lower total fat content, which is  
295 a consequence of significantly lower SFA and MUFA content. However, the amount of PUFA  
296 shows no significant difference across the six breed groups, suggesting that meat from Brahman  
297 animals would be closely aligned with the international recommendation of lean red meat to be  
298 included in a healthy balanced diet (Wyness et al., 2011; McNeill and Van Elswyk, 2012;  
299 Cashman and Hayes, 2017). There is an increasing segment of consumers interested in the taste  
300 and health benefits of products they consume (Lusk and Parker, 2009; Cashman and Hayes,  
301 2017). Recent focus of consumer interest is on weight loss and childhood obesity and the  
302 emphasis is on including protein in their diet while looking for lighter options. Leaner beef from  
303 Brahman animals could be better-fitted product for this type of consumer. In this study, steaks  
304 from Brahman steers had about half of total fat content ( $6.92 \pm 1.01$  mg/100 g meat) compared to  
305 steaks from Angus cattle ( $14.06 \pm 1.01$  mg/100 g meat) as a result of decreased SFA and MUFA  
306 concentrations, while the PUFA concentration was not significantly changed.

### 307 ***Breed Effect on Mineral Composition***

308 Least squares means of minerals concentration ( $\mu\text{g/g}$  muscle) are presented in Table 5.  
309 The mineral concentration in our study agrees with those of several other studies (Biesalski,  
310 2005; O'Neil et al., 2011) documenting the role of beef in providing essential minerals to the  
311 human diet, particularly iron, magnesium, phosphorus, potassium, and zinc. When bio-  
312 availability from other food sources is considered, the amount of iron and zinc provided through  
313 consumption of lean beef plays a critical role toward meeting the nutritional requirements of

314 these two nutrients and may provide major health benefits (Nicklas et al., 2012). The  
315 concentration of these two nutritionally critical minerals is essentially unchanged across breed  
316 groups, all having substantial nutritive value.

317 Breed effect was significant for magnesium ( $P < 0.0001$ ), phosphorus ( $P = 0.06$ ) and  
318 potassium ( $P = 0.06$ ). The concentration of these three minerals increased with the Brahman  
319 percentage in our population as indicated by a significant linear effect. The magnesium  
320 concentration in steers from the 100% Angus breed group was significantly lower compared to  
321 steers from the 50%, 75%, and 100% Brahman groups, while 75% Angus and Brangus steers  
322 had significantly different magnesium concentrations compared to 75% and 100% Brahman  
323 steers. Steers from the Brahman breed group had significantly higher phosphorus and  
324 potassium concentrations compared to 100% Angus, 75% Angus and Brangus steers. The  
325 identical direction of variation for these minerals is supported by the strong and positive  
326 genetic correlations reported between magnesium and phosphorus (0.88), magnesium and  
327 potassium (0.68), and phosphorus and potassium (0.69) (Mateescu et al., 2013). Magnesium is  
328 an essential mineral with important and extensive roles in human health including muscle and  
329 nerve function, immune system function, and bone health (Clarkson and Haymes, 1995;  
330 Genus and Bouchard, 2012; Saris et al., 2000; Spiegel, 2011; Tam et al., 2003; Orchard et al.,  
331 2014). There is an increased interest in the role of magnesium in preventing and managing  
332 disorders such as hypertension, cardiovascular disease, and diabetes (Bo and Pisu, 2008;  
333 Champagne, 2008; Houston, 2011). However, although statistically significant, the differences  
334 reported in the present study are negligible from a practical standpoint when taking into  
335 consideration the average content of these minerals and the range for their natural variation  
336 presented in this study. Irrespective of the differences found, the current study confirms

337 previous reports on the nutritionally beneficial amounts of minerals, particularly iron and zinc,  
338 in beef cattle (Mateescu et al., 2013; Ahlberg et al., 2014). Large variation in the content of  
339 these minerals was found within each breed group in the present study.

340

341

## CONCLUSION

342 One of the greatest marketing advantages of beef is that it provides a superior eating  
343 experience/taste over other protein sources. Over and above this eating experience, beef is a  
344 nutrient rich foodstuff. However, it also is perceived to have an unhealthful fatty acid  
345 composition. This study confirms that nutrient and fatty acid profiles are not uniform across  
346 cattle and variations in fatty acid composition and mineral content is partially attributable to  
347 breed composition and other genetic and management factors. As the percentage of Brahman  
348 increases, the percent of SFA out of the total fatty acids decreases and the percent of PUFA  
349 increases. These observed differences result in a more nutritionally desirable beef product,  
350 especially when we consider that the decrease in percent of SFA is mainly due to a decrease in  
351 short-chain SFA, which might have a detrimental effect on human health. Probably even more  
352 important, the relatively leaner meat of high percent Brahman steers have reduced content of  
353 SFA and MUFA but the same content of PUFA as purebred Angus, suggesting this beef product  
354 would be of interest for those consumers seeking a low fat, healthy diet.

355

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**Table 1.** Summary statistics for carcass quality, meat quality, mineral content and fatty acid composition of animals available for this study.

Trait	N	Mean	SD	Min	Max
<b>Carcass quality</b>					
BW, kg	230	34.07	6.04	18.14	50.80
WW, kg	230	238.36	40.73	120.20	353.80
HCW, kg	230	333.58	42.15	215	467.65
Dressing Percentage, %	230	60.15	2.94	53.3	70.90
Fat Thickness, cm	230	1.46	0.64	0.28	4.06
Yield Grade	230	3.43	0.80	0.60	8.10
Quality Grade	230	587.77	50.00	490.00	700.00
<b>Meat quality</b>					
LM area, cm <sup>2</sup>	230	77.81	12.09	47.74	129.03
Marbling Score <sup>1</sup>	230	416.78	89.54	280.00	700.00
WBSF, kg	230	4.52	1.18	2.10	7.90
Tenderness <sup>2</sup>	230	5.25	0.83	3.00	7.50
Juiciness <sup>2</sup>	230	4.99	0.66	3.40	6.60
Connective Tissue <sup>2</sup>	230	5.80	0.88	3.50	7.60
Flavor <sup>2</sup>	230	5.54	0.46	4.20	6.70
Off Flavor <sup>2</sup>	230	5.69	0.26	4.60	60.00
<b>Mineral content</b>					
Ca, µg/g	230	88.26	43.91	33.14	215.95
Fe, µg/g	230	14.57	3.75	4.76	27.44
K, µg/g	230	3158.86	513.79	1033.66	4658.84
Mg, µg/g	230	210.39	36.13	81.98	347.71
Na, µg/g	230	351.81	59.72	127.35	520.30
P, µg/g	230	1727.31	282.75	559.59	2796.58
Zn, µg/g	230	32.69	7.42	9.19	64.08
<b>Fatty Acid composition</b>					
SFA, %	230	48.51	3.85	38.66	64.77
MUFA, %	230	45.79	5.08	36.27	54.66
PUFA, %	230	5.70	2.58	2.69	17.39
SFA, mg/100g meat	230	4.91	3.39	0.46	15.65
MUFA, mg/100g meat	230	4.64	3.13	0.09	13.93
PUFA, mg/100g meat	230	0.49	0.25	0.05	1.85
Total Fat	230	10.04	6.71	1.34	38.26

544 <sup>1</sup>Marbling score: 100-199=Devoid, 200-299=traces, 300-399=Slight, 400-499=Small, 500-  
545 599=Modest, 600-699=Moderate, 700-799=Slightly abundant.

546 <sup>2</sup>Sensory traits: tenderness, connective tissue, beef flavor intensity, and juiciness on scales form  
547 1-8 (1=extremely tough, abundant amount, extremely bland, extremely dry; 8=extremely tender,  
548 none detected, extremely intense, extremely juicy). Off flavor was evaluated on a scale from 1-6  
549 (1=extreme off- flavor to 6=none detected).

550

**Table 2.** Least squares means and SE for carcass and meat quality characteristics in Angus (n = 39), 75% Angus (n = 33), Brangus (n = 30), 50% Angus (n = 42), 25% Angus (n = 27), and Brahman (n = 59) cattle<sup>1</sup>.

Trait	Breed Group						SE <sup>2</sup>	i <sup>3</sup>	b <sub>1</sub> <sup>3</sup>	b <sub>2</sub> <sup>3</sup>
	Angus	75% A	Brangus	50% A	25% A	Brahman				
Birth Weight, kg	33.40 <sup>bc</sup>	33.79 <sup>bc</sup>	35.99 <sup>ab</sup>	34.69 <sup>ab</sup>	37.52 <sup>a</sup>	31.51 <sup>c</sup>	0.97	32.96	3.174 <sup>S</sup>	-0.852 <sup>S</sup>
Weaning Weight, kg	254.25 <sup>a</sup>	246.88 <sup>a</sup>	251.14 <sup>a</sup>	243.65 <sup>a</sup>	238.64 <sup>a</sup>	206.68 <sup>b</sup>	5.73	268.69	4.862	-3.998 <sup>S</sup>
HCW, kg	339.61 <sup>a</sup>	349.00 <sup>a</sup>	350.48 <sup>a</sup>	338.92 <sup>a</sup>	341.30 <sup>a</sup>	304.90 <sup>b</sup>	5.48	368.09	13.654 <sup>S</sup>	-5.591 <sup>S</sup>
Dressing Percentage, %	58.88 <sup>b</sup>	60.59 <sup>a</sup>	60.90 <sup>a</sup>	60.27 <sup>a</sup>	61.23 <sup>a</sup>	60.20 <sup>a</sup>	0.38	61.242	1.443 <sup>S</sup>	-0.291 <sup>S</sup>
Fat over the eye, cm	1.64	1.42	1.45	1.4	1.44	1.4	0.11	1.676	-0.042	-
Ribeye area, cm <sup>2</sup>	79.41 <sup>a</sup>	80.77 <sup>a</sup>	81.14 <sup>a</sup>	77.07 <sup>ab</sup>	77.98 <sup>a</sup>	73.23 <sup>b</sup>	1.66	89.766	-1.769 <sup>S</sup>	-
Yield Grade	3.57	3.39	3.44	3.54	3.46	3.25	0.13	3.536	-0.066	-
Quality Grade	613.80 <sup>a</sup>	604.24 <sup>ab</sup>	612.20 <sup>a</sup>	590.82 <sup>b</sup>	582.58 <sup>b</sup>	547.37 <sup>c</sup>	7.21	619.59	0.597	-4.255 <sup>S</sup>
Marbling Score <sup>4</sup>	464.12 <sup>a</sup>	443.02 <sup>ab</sup>	456.48 <sup>ab</sup>	422.95 <sup>bc</sup>	399.01 <sup>c</sup>	352.85 <sup>d</sup>	13.31	486.02	-28.79 <sup>S</sup>	-
WBSF, kg	4.38	4.78	4.58	4.63	4.94	4.7	0.17	3.927	0.069	-
Juiciness <sup>5</sup>	5.15 <sup>a</sup>	5.12 <sup>a</sup>	5.13 <sup>a</sup>	5.04 <sup>ab</sup>	4.87 <sup>ab</sup>	4.80 <sup>b</sup>	0.11	5.049	-0.095 <sup>S</sup>	-
Flavor <sup>5</sup>	5.62	5.59	5.6	5.53	5.45	5.41	0.07	5.427	-0.057 <sup>S</sup>	-
Tenderness <sup>5</sup>	5.75 <sup>a</sup>	5.53 <sup>ab</sup>	5.51 <sup>ab</sup>	5.21 <sup>bc</sup>	5.07 <sup>cd</sup>	4.84 <sup>d</sup>	0.13	5.554	-0.231 <sup>S</sup>	-
Connective Tissue <sup>5</sup>	6.29 <sup>a</sup>	6.00 <sup>ab</sup>	6.02 <sup>ab</sup>	5.84 <sup>bc</sup>	5.49 <sup>cd</sup>	5.49 <sup>d</sup>	0.14	76.19	-0.205 <sup>S</sup>	-
Off-flavor <sup>5</sup>	5.74	5.72	5.73	5.66	5.71	5.7	0.05	5.631	-0.009	-

551 <sup>1</sup>Within each row, means without common letters differ ( $P < 0.05$ )

552 <sup>2</sup>Average SE across the breed groups

553 <sup>3</sup>Intercept (i), linear (b<sub>1</sub>) and quadratic (b<sub>2</sub>) effect of percent Brahman genetics.

554 <sup>S</sup>Linear or quadratic effect significant at  $P < 0.05$

555 <sup>4</sup>Marbling score: 100-199=Devoid, 200-299=traces, 300-399=Slight, 400-499=Small, 500-599=Modest, 600-699=Moderate, 700-799=Slightly abundant.

557 <sup>5</sup>Sensory traits: tenderness, connective tissue, beef flavor intensity, and juiciness on scales form  
558 1-8 (1=extremely tough, abundant amount, extremely bland, extremely dry; 8=extremely tender,  
559 none detected, extremely intense, extremely juicy). Off flavor was evaluated on a scale from 1-6  
560 (1=extreme off- flavor to 6=none detected).

561

**Table 3.** Least squares means and SE for fatty acid proportion (g/100 g of total fatty acids) in Angus (n = 39), 75% Angus (n = 33), Brangus (n = 30), 50% Angus (n = 42), 25% Angus (n = 27), and Brahman (n = 59) cattle<sup>1</sup>.

Trait	Breed Group						SE <sup>2</sup>	i <sup>3</sup>	b <sub>1</sub> <sup>3</sup>	b <sub>2</sub> <sup>3</sup>
	Angus	75% A	Brangus	50% A	25% A	Brahman				
C10:0	0.05 <sup>c</sup>	0.06 <sup>bc</sup>	0.06 <sup>bc</sup>	0.05 <sup>bc</sup>	0.06 <sup>b</sup>	0.07 <sup>a</sup>	0.003	0.053	-0.001	0.001 <sup>S</sup>
C11:0	0.01 <sup>b</sup>	0.02 <sup>b</sup>	0.02 <sup>b</sup>	0.02 <sup>b</sup>	0.02 <sup>b</sup>	0.04 <sup>a</sup>	0.003	0.024	-0.002	0.002 <sup>S</sup>
C12:0	0.07 <sup>c</sup>	0.08 <sup>b</sup>	0.08 <sup>b</sup>	0.07 <sup>bc</sup>	0.08 <sup>b</sup>	0.09 <sup>a</sup>	0.003	0.069	0.006 <sup>S</sup>	-
C13:0	0.01	0.02	0.02	0.01	0.01	0.02	0.002	0.014	0.001 <sup>S</sup>	-
C14:0	3.15	3.33	3.52	3.21	3.39	3.26	0.104	3.931	0.011	-
C14:1	0.70	0.82	0.87	0.81	0.83	0.86	0.046	0.833	0.029 <sup>S</sup>	-
C15:0	0.39	0.40	0.41	0.38	0.39	0.40	0.013	0.558	0.0008	-
C16:0	30.39 <sup>a</sup>	29.82 <sup>a</sup>	30.03 <sup>a</sup>	28.6 <sup>b</sup>	28.48 <sup>b</sup>	28.51 <sup>b</sup>	0.365	28.751	-0.496 <sup>S</sup>	-
C16:1	3.35 <sup>c</sup>	3.67 <sup>ab</sup>	3.73 <sup>ab</sup>	3.56 <sup>abc</sup>	3.41 <sup>bc</sup>	3.76 <sup>a</sup>	0.114	3.736	0.063	-
C17:0	1.16 <sup>a</sup>	1.12 <sup>a</sup>	1.10 <sup>a</sup>	1.10 <sup>a</sup>	1.15 <sup>a</sup>	1.14 <sup>a</sup>	0.038	1.541	-0.0008	-
C17:1	0.70 <sup>abc</sup>	0.79 <sup>ab</sup>	0.62 <sup>bc</sup>	0.61 <sup>c</sup>	0.72 <sup>abc</sup>	0.82 <sup>a</sup>	0.062	0.919	-0.112	0.033 <sup>S</sup>
C18:0	14.47	13.91	13.80	14.00	14.45	13.71	0.326	13.451	-0.115	-
C18:1	41.02	40.30	40.24	41.68	40.73	39.28	0.865	41.768	-0.359	-
C18:2n-6 <i>cis</i>	2.51 <sup>c</sup>	3.37 <sup>b</sup>	3.17 <sup>bc</sup>	3.43 <sup>b</sup>	3.73 <sup>b</sup>	4.8 <sup>a</sup>	0.270	3.298	0.530 <sup>S</sup>	-
C18:2n-6 <i>trans</i>	0.29 <sup>c</sup>	0.30 <sup>bc</sup>	0.32 <sup>ab</sup>	0.32 <sup>a</sup>	0.31 <sup>abc</sup>	0.31 <sup>ab</sup>	0.008	0.265	0.023 <sup>S</sup>	-0.005 <sup>S</sup>
C18:3n-6	0.04	0.05	0.07	0.05	0.05	0.07	0.009	0.044	0.006 <sup>S</sup>	-
C18:3n-3	0.23 <sup>c</sup>	0.25 <sup>bc</sup>	0.26 <sup>bc</sup>	0.27 <sup>b</sup>	0.27 <sup>b</sup>	0.31 <sup>a</sup>	0.008	0.202	0.019 <sup>S</sup>	-
C20:0	0.11 <sup>a</sup>	0.11 <sup>a</sup>	0.10 <sup>ab</sup>	0.11 <sup>a</sup>	0.10 <sup>ab</sup>	0.10 <sup>b</sup>	0.003	0.084	-0.003 <sup>S</sup>	-
C20:1n-9	0.28	0.30	0.29	0.03	0.27	0.28	0.010	0.319	-0.004	-
C20:2	0.08 <sup>d</sup>	0.08 <sup>cd</sup>	0.08 <sup>cd</sup>	0.09 <sup>bc</sup>	0.10 <sup>b</sup>	0.14 <sup>a</sup>	0.006	0.052	-0.0003	0.004 <sup>S</sup>
C20:3n-6	0.17 <sup>c</sup>	0.21 <sup>bc</sup>	0.23 <sup>bc</sup>	0.23 <sup>bc</sup>	0.26 <sup>b</sup>	0.34 <sup>a</sup>	0.026	0.161	0.039 <sup>S</sup>	-
C20:3n-3	0.55 <sup>c</sup>	0.67 <sup>bc</sup>	0.61 <sup>bc</sup>	0.75 <sup>bc</sup>	0.85 <sup>b</sup>	1.18 <sup>a</sup>	0.079	0.443	-0.002	0.039 <sup>S</sup>
C20:5n-3	0.05 <sup>c</sup>	0.05 <sup>bc</sup>	0.06 <sup>bc</sup>	0.06 <sup>bc</sup>	0.07 <sup>b</sup>	0.10 <sup>a</sup>	0.007	0.016	-0.001	0.004 <sup>S</sup>
C22:0	0.01	0.01	0.01	0.01	0.02	0.02	0.002	0.017	0.002 <sup>S</sup>	-
C22:2	0.01 <sup>c</sup>	0.01 <sup>bc</sup>	0.02 <sup>ab</sup>	0.02 <sup>abc</sup>	0.02 <sup>abc</sup>	0.02 <sup>a</sup>	0.003	0.034	0.002 <sup>S</sup>	-
C23:0	0.10 <sup>bc</sup>	0.13 <sup>bc</sup>	0.09 <sup>c</sup>	0.11 <sup>bc</sup>	0.14 <sup>ab</sup>	0.17 <sup>a</sup>	0.015	0.083	0.017 <sup>S</sup>	-
C22:6n-3	0.05 <sup>c</sup>	0.06 <sup>bc</sup>	0.11 <sup>ab</sup>	0.08 <sup>abc</sup>	0.05 <sup>c</sup>	0.12 <sup>a</sup>	0.020	0.009	0.012 <sup>S</sup>	-
C24:1n-9	0.05 <sup>c</sup>	0.05 <sup>c</sup>	0.10 <sup>ab</sup>	0.06 <sup>bc</sup>	0.04 <sup>c</sup>	0.10 <sup>a</sup>	0.014	0.025	0.009 <sup>S</sup>	-
SFA	49.92 <sup>a</sup>	49 <sup>abc</sup>	49.23 <sup>ab</sup>	47.69 <sup>bc</sup>	48.29 <sup>abc</sup>	47.52 <sup>c</sup>	0.602	48.031	-0.566 <sup>S</sup>	-
MUFA	46.10	45.93	45.84	47.02	45.99	45.10	0.830	47.531	-0.235	-
PUFA	3.98 <sup>c</sup>	5.06 <sup>bc</sup>	4.93 <sup>bc</sup>	5.29 <sup>b</sup>	5.72 <sup>b</sup>	7.38 <sup>a</sup>	0.391	4.438	0.802 <sup>S</sup>	-

<sup>1</sup>Within each row, means without common letters differ ( $P < 0.05$ )

<sup>2</sup>Average SE across the breed groups

<sup>3</sup>Intercept (i), linear (b<sub>1</sub>) and quadratic (b<sub>2</sub>) effect of percent Brahman genetics.

<sup>S</sup>Linear or quadratic effect significant at  $P < 0.05$

**Table 4.** Least squares means and SE for fatty acid concentration (mg/100 g meat) in Angus (n = 39), 75% Angus (n = 33), Brangus (n = 30), 50% Angus (n = 42), 25% Angus (n = 27), and Brahman (n = 59) cattle<sup>1</sup>.

Trait	Breed Group						SE <sup>2</sup>	i <sup>3</sup>	b <sub>1</sub> <sup>3</sup>	b <sub>2</sub> <sup>3</sup>
	Angus	75% A	Brangus	50% A	25% A	Brahman				
C10:0	0.007 <sup>a</sup>	0.006 <sup>ab</sup>	0.006 <sup>ab</sup>	0.006 <sup>ab</sup>	0.006 <sup>ab</sup>	0.005 <sup>b</sup>	0.0012	0.005	-0.0006 <sup>S</sup>	-
C11:0	0.002	0.002	0.001	0.002	0.001	0.002	0.0002	0.001	5.68x10 <sup>-6</sup>	-
C12:0	0.009	0.008	0.009	0.008	0.008	0.006	0.0011	0.007	-0.0007 <sup>S</sup>	-
C13:0	0.001	0.002	0.002	0.002	0.001	0.001	0.0002	0.001	-0.0001 <sup>S</sup>	-
C14:0	0.446 <sup>a</sup>	0.380 <sup>a</sup>	0.401 <sup>a</sup>	0.352 <sup>a</sup>	0.336 <sup>a</sup>	0.230 <sup>b</sup>	0.0672	0.354	-0.052 <sup>S</sup>	-
C14:1	0.103 <sup>a</sup>	0.095 <sup>a</sup>	0.096 <sup>a</sup>	0.090 <sup>a</sup>	0.079 <sup>ab</sup>	0.056 <sup>b</sup>	0.0168	0.086	-0.012 <sup>S</sup>	-
C15:0	0.054 <sup>a</sup>	0.044 <sup>ab</sup>	0.047 <sup>ab</sup>	0.040 <sup>b</sup>	0.034 <sup>bc</sup>	0.025 <sup>c</sup>	0.0054	0.055	-0.007 <sup>S</sup>	-
C16:0	4.301 <sup>a</sup>	3.468 <sup>ab</sup>	3.481 <sup>ab</sup>	3.129 <sup>b</sup>	2.814 <sup>bc</sup>	1.967 <sup>c</sup>	0.6962	3.111	-0.556 <sup>S</sup>	-
C16:1	0.475 <sup>a</sup>	0.421 <sup>ab</sup>	0.417 <sup>ab</sup>	0.391 <sup>ab</sup>	0.321 <sup>bc</sup>	0.250 <sup>c</sup>	0.0682	0.387	-0.058 <sup>S</sup>	-
C17:0	0.156 <sup>a</sup>	0.123 <sup>b</sup>	0.126 <sup>ab</sup>	0.115 <sup>b</sup>	0.101 <sup>bc</sup>	0.073 <sup>c</sup>	0.0147	0.154	-0.019 <sup>S</sup>	-
C17:1	0.095 <sup>a</sup>	0.089 <sup>ab</sup>	0.073 <sup>ab</sup>	0.068 <sup>bc</sup>	0.063 <sup>bc</sup>	0.048 <sup>c</sup>	0.0121	0.086	-0.012 <sup>S</sup>	-
C18:0	2.015 <sup>a</sup>	1.612 <sup>ab</sup>	1.592 <sup>b</sup>	1.519 <sup>b</sup>	1.443 <sup>b</sup>	0.955 <sup>c</sup>	0.3157	1.450	-0.243 <sup>S</sup>	-
C18:1	5.695 <sup>a</sup>	4.789 <sup>ab</sup>	4.629 <sup>ab</sup>	4.461 <sup>b</sup>	3.949 <sup>bc</sup>	2.843 <sup>c</sup>	0.8208	4.516	-0.686 <sup>S</sup>	-
C18:2n-6 <i>cis</i>	0.346	0.337	0.337	0.335	0.312	0.272	0.0404	0.296	-0.019 <sup>S</sup>	-
C18:2n-6 <i>trans</i>	0.043 <sup>a</sup>	0.037 <sup>ab</sup>	0.037 <sup>ab</sup>	0.036 <sup>ab</sup>	0.031 <sup>bc</sup>	0.021 <sup>c</sup>	0.0085	0.030	-0.005 <sup>S</sup>	-
C18:3n-6	0.007	0.005	0.006	0.005	0.005	0.004	0.0014	0.004	-0.0006 <sup>S</sup>	-
C18:3n-3	0.034 <sup>a</sup>	0.029 <sup>ab</sup>	0.029 <sup>ab</sup>	0.029 <sup>ab</sup>	0.026 <sup>bc</sup>	0.021 <sup>c</sup>	0.0063	0.023	-0.003 <sup>S</sup>	-
C20:0	0.015 <sup>a</sup>	0.013 <sup>ab</sup>	0.012 <sup>ab</sup>	0.012 <sup>b</sup>	0.011 <sup>b</sup>	0.007 <sup>c</sup>	0.0031	0.009	-0.002 <sup>S</sup>	-
C20:1n-9	0.040 <sup>a</sup>	0.037 <sup>a</sup>	0.034 <sup>ab</sup>	0.032 <sup>ab</sup>	0.027 <sup>bc</sup>	0.019 <sup>c</sup>	0.0061	0.034	-0.005 <sup>S</sup>	-
C20:2	0.010	0.009	0.009	0.010	0.009	0.009	0.0008	0.025	-0.0004	-
C20:3n-6	0.022	0.021	0.023	0.022	0.022	0.018	0.0044	0.014	-0.0007	-
C20:3n-3	0.071	0.068	0.060	0.075	0.071	0.066	0.0195	0.032	-0.0005	-
C20:5n-3	0.006	0.006	0.005	0.007	0.006	0.006	0.0024	0.001	0.00006	-
C22:0	0.001	0.001	0.001	0.001	0.001	0.001	0.0002	0.001	-0.0001 <sup>S</sup>	-
C22:2	0.001 <sup>b</sup>	0.001 <sup>bc</sup>	0.002 <sup>a</sup>	0.001 <sup>b</sup>	0.001 <sup>bc</sup>	0.001 <sup>c</sup>	0.0001	0.002	0.0003 <sup>S</sup>	-0.0001 <sup>S</sup>
C23:0	0.013	0.012	0.010	0.010	0.012	0.010	0.0027	0.007	-0.0006	-
C22:6n-3	0.006	0.006	0.008	0.007	0.006	0.007	0.0019	0.001	0.0002	-
C24:1n-9	0.006	0.005	0.007	0.005	0.004	0.006	0.0029	0.002	-0.0001	-
SFA	7.023 <sup>a</sup>	5.670 <sup>ab</sup>	5.688 <sup>ab</sup>	5.195 <sup>b</sup>	4.771 <sup>b</sup>	3.282 <sup>c</sup>	1.0781	5.157	-0.885 <sup>S</sup>	-
MUFA	6.414 <sup>a</sup>	5.437 <sup>ab</sup>	5.256 <sup>ab</sup>	5.048 <sup>b</sup>	4.443 <sup>bc</sup>	3.221 <sup>c</sup>	0.9245	5.111	-0.774 <sup>S</sup>	-
PUFA	0.547	0.520	0.516	0.527	0.490	0.426	0.0876	0.411	-0.030 <sup>S</sup>	-
Total Fat	13.985 <sup>a</sup>	11.627 <sup>ab</sup>	11.461 <sup>ab</sup>	10.770 <sup>b</sup>	9.705 <sup>bc</sup>	6.929 <sup>c</sup>	2.088	10.679	-1.686 <sup>S</sup>	-

<sup>1</sup>Within each row, means without common letters differ ( $P < 0.05$ )

<sup>2</sup>Average SE across the breed groups

<sup>3</sup>Intercept (i), linear (b<sub>1</sub>) and quadratic (b<sub>2</sub>) effect of percent Brahman genetics.

<sup>S</sup>Linear or quadratic effect significant at  $P < 0.05$

**Table 5.** Least squares means and SE for minerals concentration ( $\mu\text{g/g}$  muscle) in Angus ( $n = 39$ ), 75% Angus ( $n = 33$ ), Brangus ( $n = 30$ ), 50% Angus ( $n = 42$ ), 25% Angus ( $n = 27$ ), and Brahman ( $n = 59$ ) cattle<sup>1</sup>.

Trait	Breed Group						SE <sup>2</sup>	i <sup>3</sup>	b <sub>1</sub> <sup>3</sup>
	Angus	75% A	Brangus	50% A	25% A	Brahman			
Iron	14.58	14.44	13.55	15.28	14.70	14.62	0.65	14.56	0.05
Magnesium	196.83 <sup>c</sup>	198.88 <sup>bc</sup>	198.94 <sup>bc</sup>	212.79 <sup>ab</sup>	216.42 <sup>ab</sup>	222.86 <sup>a</sup>	8.18	202.97	7.40 <sup>S</sup>
Zinc	33.79	32.81	31.871	33.23	34.531	31.859	1.48	34.22	-0.30
Sodium	338.27	348.38	339.68	362.11	356.30	360.40	10.56	339.02	4.99
Phosphorus	1651.80 <sup>b</sup>	1677.07 <sup>b</sup>	1655.55 <sup>b</sup>	1709.01 <sup>ab</sup>	1748.20 <sup>ab</sup>	1803.98 <sup>a</sup>	58.06	1743.39	41.45 <sup>S</sup>
Potassium	3016.29 <sup>b</sup>	3041.87 <sup>b</sup>	3015.07 <sup>b</sup>	3138.67 <sup>ab</sup>	3221.53 <sup>ab</sup>	3273.92 <sup>a</sup>	103.75	3157.68	74.50 <sup>S</sup>

<sup>1</sup>Within each row, means without common letters differ ( $P < 0.05$ )

<sup>2</sup>Average SE across the breed groups

<sup>3</sup>Intercept (i) and linear (b<sub>1</sub>) effect of percent Brahman genetics. No quadratic (b<sub>2</sub>) effect was significant.

<sup>S</sup>Linear effect significant at  $P < 0.05$ .