

# Crossbreeding applied to systems of beef cattle production to improve performance traits and carcass quality

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Crossbreeding represents an important technique to improve growth, beef quality and adaptability in beef production systems in tropical countries. The aim of this study was to evaluate sire and dam breed effects on performance and carcass traits of crossbred cattle produced in a tropical environment. Heifers and steers were evaluated during the pre-weaning, the post-weaning (n = 173) and the finishing phase (n = 123). Animals were produced by mating Nellore (N N). Angus  $\times$  Nellore (A N) and Caracu  $\times$  Nellore (**C**\_**N**) dams with Braford, Charbray and Caracu sires. After weaning, animals were raised grazing on Marandu grass for 12 months; thereafter they were housed in individual pens and finished in a feedlot, receiving a total mixed ration. Ultrasound carcass evaluations were performed to determine ribeye area ( $R_A$ ), backfat thickness ( $B_T$ ) and rump fat thickness (**R** T). A N progeny were heavier at birth than N N (P < 0.05), and Braford progeny had greater birth BW than Caracu (P < 0.05). Greater weaning BW was observed in the A\_N and C\_N offspring compared to N\_N (P < 0.01). Greater average daily gain during the post-weaning period was verified in the N\_N progeny compared to C\_N (P < 0.05). No dam or sire breed effects were observed for BW at the end of the post-weaning period (P > 0.05). Progeny of N N cows had greater B T (P < 0.05) and  $R_T$  (P < 0.01) at the end of the post-weaning period in relation to C\_N. Greater R\_A was observed in the Caracu progeny than in the Braford (P < 0.05), which showed greater R T than the Charbray progeny at the end of the post-weaning period (P < 0.05). No dam or sire breed effects were verified for final BW at the feedlot or for feed efficiency traits (P > 0.05). A N progeny were superior in final B\_T compared to  $C_N$  (P < 0.01), and Braford progeny had greater R\_T at the end of finishing than Charbray (P = 0.05). The use of crossbred dams allows an increase in productivity until weaning, but this is not maintained in the post-weaning and finishing periods. The use of Braford sires produces similar growth performance in the different stages of the production system to those seen with Charbray and Caracu sires but generates animals with higher fat thickness at the end of finishing, which may improve carcass guality and commercial value.

Keywords: adapted breeds, backfat, Bos indicus, crossbreeding, growth

# Implications

Beef industries in tropical regions that rely on pasture-based production systems may benefit from crossbreeding schemes using zebu, adapted taurine, composites and British breeds, due to complementarity of characteristics like improved adaptability, greater resistance against parasites, higher growth rate and superior meat quality. Recommendations on the use of Nellore (zebu), Caracu (tropically adapted Brazilian native taurine), Charbray (Charolais × zebu composite),

Braford (British  $\times$  zebu composite) and Angus (British) may be provided, so improvements in growth performance and in carcass quality can take place. Consequently, greater revenues from enhanced productivity and higher carcass commercial value can be achieved.

# Introduction

Success in beef production depends on the effects of genetic and environmental (e.g. nutrition, health and climate) factors, as well as the interaction among these elements. It is known that approximately 80% of the meat produced in Brazil, one of

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the largest producers in the world, originates from zebu breeds, predominantly the Nellore breed (Ferraz and Felício, 2010). Zebu breeds stand out for their greater adaptability to the climatic conditions present in the tropics and greater resistance to ectoparasites and certain diseases; however, they usually show lower growth rates and worse meat quality, especially with regard to tenderness, when compared to taurine breeds (Wheeler *et al.*, 2010; Elzo *et al.*, 2012). In this context, crossbreeding between *Bos indicus* and *Bos taurus* animals is an effective strategy to optimise the use of genetic and environmental resources, since it allows genetic gains through heterosis (Gama *et al.*, 2013) and a combination of desirable traits through breed complementarity, promoting an increase in the productive potential of animals in tropical environments (Miguel *et al.*, 2014).

However, it is important to note that increases in the proportion of genes from zebu animals in the crossbred progeny result in lower reproductive precocity, lower deposition of fat in the carcass and less tender beef (Rotta *et al.*, 2009). The challenge of maintaining both adaptation to tropical conditions and sufficient taurine genes may be overcome by using crossbreeding schemes with Nellore or crossbred dams (*Bos taurus* × *Bos indicus*), and sires of composite breeds (*Bos taurus* × *Bos indicus*), such as Braford and Charbray, or sires of taurine breeds (*Bos taurus*) adapted to tropical conditions, such as the Brazilian native Caracu breed. An added advantage of this approach is that it allows for the possibility of using natural mating.

Despite the advantages of crossbreeding in terms of carcass and meat quality and performance, it is necessary to identify crossbreeding schemes that are better suited to particular environments and production systems, such as systems developed in tropical environments that are designed for the slaughter of young animals. Thus, the objective of this study was to evaluate the effects of different sire and dam breeds, adapted to the tropical climate, on the performance and carcass traits assessed by ultrasound of crossbred cattle reared in tropical pasture and finished in feedlot.

# Materials and methods

This study was carried out at Embrapa Beef Cattle in Campo Grande, Brazil (20°27′ S and 54°37′ W, at 530 m above sea level). The climate, according to the Köppen climate classification, is type AW (rainy tropical savanna) with a defined dry season from May to September. Mean annual temperature and precipitation are 23.4°C and 1449 mm, respectively.

Experimental animals were generated from mating Nellore (**N**\_**N**),  $\frac{1}{2}$  Angus +  $\frac{1}{2}$  Nellore (**A**\_**N**) and  $\frac{1}{2}$  Caracu +  $\frac{1}{2}$  Nellore (**C**\_**N**) cows (4 to 8 years old) with Braford, Charbray and Caracu sires (9, 10 and 18 sires, respectively) using fixed-time artificial insemination in breeding seasons carried out from November 2012 to February 2013 (batch 1) and November 2013 to February 2014 (batch 2). Crossbred progenies were evaluated along the pre-weaning (0 to 8 months of age, n = 173), post-weaning

(8 to 20 months of age, n = 173) and finishing (20 to 23 months of age, n = 119) phases. Either for producing the cows or their progenies, sires were chosen from animal breeding programmes for their respective breeds, opting for individuals from different lineages so that each breed could be widely represented. Table 1 summarises the different phases of the study and the number of observations.

During the pre-weaning phase, cows and calves were maintained in *Brachiaria* spp. and *Panicum* spp. pastures receiving mineral supplements. Calf BW was obtained at birth and at weaning (8 months old on average). Weaning BW was adjusted for day 240 regressing calf BW on time. Pre-weaning phase average daily weight gain (**ADG**) was calculated for calves from the difference between birth and weaning BW.

Following weaning, calves were enrolled in the postweaning phase, which was split into two consecutive phases: phase 1 (**P1**, June to October) and phase 2 (**P2**, November to May). This was carried out to consider the expected effects of forage quality seasonality on growth performance. Lower precipitation and temperatures from June to November (P1) result in low quality pastures and low ADG, while higher precipitation and temperatures from December to May (P2) lead to higher quality pastures and greater ADG. In this phase, cattle were allotted to one of eight paddocks (single sex and balanced for the different breeds) of 8 ha each, planted with *Brachiaria brizantha* cv. Marandu, at continuous stocking with a fixed stocking rate.

Cattle had access to protein supplements (1 g/kg BW per day) in phase 1 and to mineral supplements (*ad libitum*) in phase 2. Animals were treated ectopically to control ticks and horn flies (cypermethrin pour-on topical solution) and endoparasites (injectable doramectin 1%). Male calves were castrated at 14 months of age and cattle were weighed following 16-h fasting every 56 days, on average. Average daily weight gain was calculated separately for P1, P2 and the total post-weaning period (P1 + P2).

Pasture sampling was conducted five times a year, distributed through the different seasons. Leaf samples were ground to 1 mm and analysed for percentages of CP, NDF and ADL, using NIRS, as described by Marten *et al.* (1989). The forage mass, green forage allowance and stocking rate varied from 10.5 to 16.1 tons DM/ha, 11.4 to 18.5 kg DM/kg BW and 0.76 to 1.30 animal units/ha in the 2 years, respectively. Crude protein, NDF and ADL contents varied from 6.6 to 8.7%, 66.9 to 73.7% and 2.1 to 2.9%, respectively.

On the final weighing of the post-weaning phase, cattle were scanned using a real-time ESAOTE Piemedical Aquila Vet (ESAOTE Pie Medical Imaging, Inc., Maastricht, The Netherlands) ultrasound unit equipped with an 18-cm, 3.5-MHz linear array transducer and a coupled acoustic guide. Vegetable oil was used as an acoustic couplant. Ultrasound images of the longissimus between the 12th and 13th ribs were used to measure ribeye area (**R\_A**) and subcutaneous backfat thickness (**B\_T**), and an image of biceps femoris muscle was used to obtain rump fat thickness (**R\_T**). Ultrasound

| Phase   | Batch 1                         | Batch 2                        |
|---|---------------------------------|--------------------------------|
| Breeding season   | November 2011 to February 2012  | November 2012 to February 2013 |
| Nutrition: pasture + mineral supplement                   | Cows                            | Cows:                          |
|   | Nellore: 38                     | Nellore: 45                    |
|   | 1/2 Angus + 1/2 Nellore: 24     | 1/2 Angus + 1/2 Nellore: 24    |
|   | 1/2 Caracu + 1/2 Nellore: 46    | 1/2 Caracu + 1/2 Nellore: 45   |
|   | Sires                           | Sires                          |
|   | Caracu: 18                      | Caracu: 18                     |
|   | Charbray: 10                    | Charbray: 10                   |
|   | Braford: 9                      | Braford: 9                     |
| Calving season<br>Nutrition: pasture + mineral supplement | August 2012 to December 2012    | August 2013 to November 2013   |
| Pre-weaning   | August 2012 to June 2013        | August 2013 to June 2014       |
| Nutrition: pasture + mineral supplement                   | Age: birth to $\sim$ 8 m        | Age: birth to $\sim$ 8 m       |
|   | Female: 46                      | Female: 41                     |
|   | Male: 42                        | Male: 44                       |
|   | Total: 88                       | Total: 85                      |
| Post-weaning 1 (P1)                                       | June 2013 to November 2013      | June 2014 to November 2014     |
| Nutrition: pasture + protein supplement (dry season)      | Age: $\sim$ 8 m to $\sim$ 13 m  | Age: $\sim$ 8 m to $\sim$ 13 m |
|   | Female: 46                      | Female: 41                     |
|   | Male: 42                        | Male: 44                       |
|   | Total: 88                       | Total: 85                      |
| Post-weaning 2 (P2)                                       | November 2013 to June 2014      | November 2013 to June 2014     |
| Nutrition: pasture + protein-energy                       | Age: $\sim$ 13 m to $\sim$ 20 m | Age: ~13 to ~20 m              |
| supplement (rainy season)                                 | Female: 46                      | Female: 41                     |
|   | Male: 42                        | Male: 44                       |
|   | Total: 88                       | Total: 85                      |
| Finishing Nutrition: Feedlot                              | June 2014 to September 2014     | June 2015 to September 2015    |
|   | Age: ~20 m to ~23 m             | Age: ~20 m to ~23 m            |
|   | Female: 26                      | Female: 34                     |
|   | Male: 29                        | Male: 34                       |
|   | Total: 55                       | Total: 68                      |

 Table 1 Summary of experimental phases and number of animals<sup>1</sup>

m = month.

<sup>1</sup> Cattle were generated from mating Nellore, ½ Angus + ½ Nellore and ½ Caracu + ½ Nellore cows with Braford, Charbray and Caracu sires.

images were interpreted using Lince software (M&S Consultoria Agropecuária Ltda, Pirassununga, SP, Brazil).

Following the post-weaning phase, cattle were housed in individual pens that were  $2 \times 19$  m and soil-surfaced. Cattle were allowed to adapt to the diet and feeding procedures for 24 days. After the adaptation period, the experimental diet was offered *ad libitum*, at 0800 h and 1500 h, so that 5% refusals were allowed. The duration of the feeding period was 83 and 84 days for heifers and 119 and 98 days for steers from batches 1 and 2, respectively.

Diet consisted of 35% sorghum silage, 49.8% ground dry corn, 9% soybean hulls, 3.6% soybean meal, 1.1% urea and 1.5% mineral premix, in dry basis, and the same composition was used in both batches. Predicted metabolisable energy (**ME**) content was 2.66 Mcal ME/kg DM, according to National Research Council (NRC, 1996). Crude protein and NDF contents were 12% and 38%, respectively, and were determined according to the Association of Official Analytical Chemists (AOAC, 1990) and Van Soest *et al.* (1991). Feed refusals were weighed daily before the morning feed delivery to calculate the daily dry matter intake (**DMI**). Cattle were weighed following 16 h of feed withdrawal every 28 days on average. Average daily gain was computed as the slope of the linear regression of BW on feeding days. Gross feed efficiency (ADG/DMI) and feed conversion ratio (DMI/ADG) were calculated. Cattle were also scanned for carcass traits at the beginning and the end of the feeding period, according to the procedures described previously.

The data were analysed for outliers (> 3 standard deviations), homogeneity of variance (Hartley test) and normality of residuals (Shapiro–Wilk test). Linear mixed models were fitted in restricted maximum likelihood (**REML**) analysis to evaluate the effects of breed, using PROC MIXED of SAS (version 9.2, SAS Institute Inc., Cary, NC, USA).

For traits evaluated in the pre-weaning phase, the model included as fixed effects: (i) contemporary group (eight levels); (ii) sire breed (three levels – Braford, Caracu and Charbray); (iii) dam breed (three levels –  $A_N$ ,  $C_N$  and  $N_N$ ; (iv) the interaction between sire and dam breed; (v) cow age at calving (linear and quadratic covariates); and (vi) and the deviation of calf birth date from the average birth date of the respective birth season, nested in the

|                       |                                      | Dam breed                 |                           |         |                           | Sire breed                 |                           |         |
|-----------------------|--------------------------------------|---------------------------|---------------------------|---------|---------------------------|----------------------------|---------------------------|---------|
| Variables             | A_N                                  | C_N                       | N_N                       | P-value | Braford                   | Charbray                   | Caracu                    | P-value |
| Pre-weaning period    |                                      |                           |                           |         |                           |                            |                           |         |
| BW at birth (kg)      | 34.0 <sup>a</sup> (0.8) <sup>2</sup> | 32.0 <sup>ab</sup> (0.6)  | 31.5 <sup>b</sup> (0.8)   | 0.042   | 33.5 <sup>a</sup> (1.0)   | 33.0 <sup>ab</sup> (1.0)   | 31.0 <sup>b</sup> (0.5)   | 0.049   |
| BW at weaning (kg)    | 228 <sup>a</sup> (4.7)               | 225 <sup>a</sup> (3.7)    | 208b (4.7)                | 0.0008  | 225 (5.3)                 | 219 (5.6)                  | 217 (2.8)                 | 0.391   |
| ADG (kg/day)          | 0.809 <sup>a</sup> (0.02)            | 0.808 <sup>a</sup> (0.02) | 0.727 <sup>b</sup> (0.02) | 0.0003  | 0.803 (0.02)              | 0.765 (0.02)               | 0.776 (0.01)              | 0.408   |
| Post-weaning period   |                                      |                           |                           |         |                           |                            |                           |         |
| Final BW, P1 (kg)     | 258 (5.1)                            | 258 (4.1)                 | 253 (4.9)                 | 0.658   | 257 (5.9)                 | 255 (6.1)                  | 257 (3.2)                 | 0.958   |
| Final BW, P2 (kg)     | 395 (7.4)                            | 394 (6.1)                 | 397 (7.7)                 | 0.918   | 394 (8.3)                 | 399 (8.5)                  | 393 (5.0)                 | 0.796   |
| ADG, P1 (kg/day)      | 0.396 <sup>ab</sup> (0.02)           | 0.368 <sup>b</sup> (0.02) | 0.432 <sup>a</sup> (0.02) | 0.014   | 0.371 <sup>b</sup> (0.03) | 0.401 <sup>ab</sup> (0.03) | 0.423 <sup>a</sup> (0.02) | 0.041   |
| ADG, P2 (kg/day)      | 0.661 (0.02)                         | 0.657 (0.02)              | 0.691 (0.02)              | 0.313   | 0.667 (0.02)              | 0.680 (0.02)               | 0.661 (0.01)              | 0.742   |
| ADG, P1 + P2 (kg/day) | 0.565 <sup>ab</sup> (0.02)           | 0.549 <sup>b</sup> (0.02) | 0.596 <sup>a</sup> (0.02) | 0.035   | 0.555 (0.02)              | 0.579 (0.02)               | 0.575 (0.01)              | 0.421   |

Table 2 Effect of dam and sire breeds on progeny growth performance during the pre- and post-weaning periods<sup>1</sup>

A\_N = 1/2 Angus + 1/2 Nellore cows, C\_N = 1/2 Caracu + 1/2 Nellore cows; N\_N = Nellore cows; P1 = phase 1 of post-weaning period; P2 = phase 2 of post-weaning period; ADG = average daily gain.

<sup>1</sup> Cattle were generated from mating Nellore, ½ Angus + ½ Nellore and ½ Caracu + ½ Nellore cows with Braford, Charbray and Caracu sires.

<sup>2</sup> Standard error of mean.

 $^{a,b}$  Values within a row with different superscripts differ significantly at P < 0.05.

contemporary group (linear covariate). Contemporary groups were composed of batch (two levels – batches 1 and 2), calving season (two levels – season 1 = August to September, and season 2 = October to November) and sex (two levels – male and female). To accommodate the effect of sampling some few sires within a breed, random effect of individual sire nested within sire breed was included in the analysis model. As covariance structure, independent covariance model was adopted, where the within-subject error correlation is zero.

To analyse the variables obtained in the post-weaning phase, a similar model to the one adopted to pre-weaning phase was employed. Cattle age at the end of phase 1 was added as a linear covariate and paddock as a random effect. For variables obtained in the finishing phase, the model included as fixed effects: (i) contemporary group (as previously described); (ii) sire breed (as previously described); (iii) dam breed (as previously described); (iv) the interaction sire by dam breed; (v) cattle age at the beginning of the finishing period (linear covariate); and (vi) days on feed (linear covariate). Individual sire nested within sire breed and paddock were included as random effects. As covariance structure, independent covariance model was adopted, where the within-subject error correlation is zero. Additionally, contrasts were performed to compare: (1) progeny of crossbred cows (A\_N and C\_N) v. progeny of purebred N\_N cows and (2) composite breed sire progeny (Braford and Charbray) v. Caracu sire progeny. When needed, means were compared by the Tukey-Kramer adjusted test. A treatment difference of  $P \le 0.05$  was considered significant, and 0.05 < P < 0.10 was considered a tendency.

# Results

Table 2 shows the results of growth performance of the progenies during the pre- and post-weaning periods. There was no interaction between dam and sire breed genetic effect for birth BW, weaning BW, pre-weaning phase ADG, final BW in post-weaning P1, final BW in post-weaning P2, post-weaning P2 ADG or post-weaning P1 + P2 ADG (P > 0.05). By comparing dam breeds, calves from A\_N cows were heavier than those from N\_N cows (P < 0.05). Calves from C\_N cows did not differ from the other two breeds (P > 0.05). Calves from A\_N and C\_N had greater weaning weights than calves from N\_N cows (P < 0.01) but did not differ from each other (P > 0.05). This result is consistent with the verified ADG, as calves from A\_N and C\_N cows (P < 0.01).

Sire breed genetic effects were observed for birth BW (P < 0.05). Calves from Braford sires were heavier at birth than calves from Caracu sires, whereas calves from Charbray calves were intermediate. Sire breed did not affect weaning BW or pre-weaning ADG (P > 0.05). For the post-weaning period, no sire or dam breed effects were observed for P1 final BW, P2 final BW or P2 ADG (P > 0.05). Calves from N\_N cows had greater P1 and P1 + P2 ADG when compared to C\_N cows (P < 0.01), and A\_N cows were intermediate (P > 0.05). Progeny of Caracu sires had higher P1 ADG than the progeny of Braford sires (P < 0.05), but neither breeds differed from Charbray (P > 0.05).

There was a significant interaction effect of dam and sire breeds on P1 ADG (P < 0.05, data not shown). Progeny that were  $\frac{1}{2}$  Caracu +  $\frac{1}{2}$  Nellore had greater ADG than all other breeds. Braford × C\_N had the lowest ADG, which did not differ from Charbray × A\_N. Braford × A\_N, Braford × N\_N, Charbray × C\_N, Charbray × N\_N, Caracu × A\_N and Caracu × C\_N did not differ from each other.

Table 3 shows the results of ultrasound carcass traits of the progenies, during the pre- and post-weaning periods. There were no interaction effects of dam and sire breeds on ultrasound carcass traits, measured at the end of the post-weaning P2 (P > 0.05). There were no effects of dam breed (P > 0.05) for R\_A, but progeny of N\_N cows had higher B\_T and R\_T (P < 0.01) when compared to progeny of

|                        |                           | 1 5                      | ,                        |         | 1                         | 51                         |                           |         |
|------------------------|---------------------------|--------------------------|--------------------------|---------|---------------------------|----------------------------|---------------------------|---------|
|                        |                           | Dam breed                |                          |         |                           | Sire breed                 |                           |         |
| Variables              | A_N                       | C_N                      | N_N                      | P-value | Braford                   | Charbray                   | Caracu                    | P-value |
| R_A (cm <sup>2</sup> ) | 67.68 (1.96) <sup>2</sup> | 69.40 (1.63)             | 68.53 (2.03)             | 0.704   | 65.26 <sup>b</sup> (2.15) | 69.68 <sup>ab</sup> (2.21) | 70.67ª (1.33)             | 0.042   |
| B_T (mm)               | 3.68 <sup>ab</sup> (0.32) | 3.13 <sup>b</sup> (0.27) | 4.02 <sup>a</sup> (0.35) | 0.046   | 3.71 (0.36)               | 3.35 (0.38)                | 3.77 (0.21)               | 0.572   |
| R_T (mm)               | 5.17 <sup>ab</sup> (0.43) | 4.41 <sup>b</sup> (0.37) | 5.66 <sup>a</sup> (0.44) | 0.014   | 5.65 <sup>a</sup> (0.51)  | 4.42 <sup>b</sup> (0.52)   | 5.17 <sup>ab</sup> (0.32) | 0.042   |

 Table 3 Effect of dam and sire breeds on progeny ultrasound carcass traits at the end of post-weaning period<sup>1</sup>

 $A_N = \frac{1}{2}$  Angus +  $\frac{1}{2}$  Nellore cows,  $C_N = \frac{1}{2}$  Caracu +  $\frac{1}{2}$  Nellore cows;  $N_N =$  Nellore cows;  $R_A =$  ultrasound ribeye area;  $B_T =$  ultrasound backfat thickness;  $R_T =$  ultrasound rump fat thickness.

<sup>1</sup> Cattle were generated from mating Nellore, <sup>1</sup>/<sub>2</sub> Angus + <sup>1</sup>/<sub>2</sub> Nellore and <sup>1</sup>/<sub>2</sub> Caracu + <sup>1</sup>/<sub>2</sub> Nellore cows with Braford, Charbray and Caracu sires.

<sup>2</sup> Standard error of mean.

 $^{a,b}$  Values within a row with different superscripts differ significantly at P < 0.05.

Table 4 Effect of dam breed on progeny growth performance and ultrasound carcass traits in the pre- and post-weaning periods<sup>1</sup>

|   | Dam ł                    |              |         |
|---|--------------------------|--------------|---------|
| Variables                                 | Crossbred <sup>2</sup>   | Nellore      | P-value |
| Pre-weaning period                        |                          |              |         |
| BW at birth (kg)                          | 33.0 (0.74) <sup>3</sup> | 31.5 (0.81)  | 0.058   |
| BW at weaning (kg)                        | 227 (4.2)                | 208 (4.7)    | 0.0002  |
| Average daily gain (kg/day)               | 0.809 (0.02)             | 0.727 (0.02) | <0.0001 |
| Post-weaning period                       |                          |              |         |
| Final BW, P1 (kg)                         | 258 (4.6)                | 253 (4.9)    | 0.363   |
| Final BW, P2 (kg)                         | 394 (6.7)                | 397 (7.7)    | 0.722   |
| Average daily gain, P1 (kg/day)           | 0.382 (0.02)             | 0.432 (0.02) | 0.009   |
| Average daily gain, P2 (kg/day)           | 0.659 (0.02)             | 0.691 (0.02) | 0.131   |
| Average daily gain, P1 + P2 (kg/day)      | 0.557 (0.02)             | 0.596 (0.02) | 0.017   |
| Ultrasound ribeye area (cm <sup>2</sup> ) | 68.54 (1.80)             | 68.53 (2.03) | 0.995   |
| Ultrasound backfat thickness (mm)         | 3.41 (0.30)              | 4.02 (0.35)  | 0.082   |
| Ultrasound rump fat thickness (mm)        | 4.80 (0.40)              | 5.66 (0.44)  | 0.026   |

P1 = post-weaning phase 1; P2 = post-weaning phase 2.

<sup>1</sup> Cattle were generated from mating Nellore, <sup>1</sup>/<sub>2</sub> Angus + <sup>1</sup>/<sub>2</sub> Nellore and <sup>1</sup>/<sub>2</sub> Caracu + <sup>1</sup>/<sub>2</sub> Nellore cows with Braford, Charbray and Caracu sires.

<sup>2</sup> ½ Caracu + ½ Nellore and ½ Angus + ½ Nellore cows.

<sup>3</sup> Standard error of mean.

C\_N cows, whereas A\_N progeny were intermediate (P > 0.05). Progeny of Caracu sires had greater R\_A than did those of Braford sires, but did not differ from values for Charbray progeny, which were similar to values for Braford progeny as well (P > 0.05). No sire breed effects on B\_T were observed (P > 0.05), but progeny from Braford sires had greater R\_T when compared to those of Charbray sires (P < 0.05), whereas the values for Caracu were similar to those of the other two breeds (P > 0.05).

Table 4 shows the results from the contrasts approach to compare progenies from crossbred (A\_N and C\_N) and N\_N cows for growth and carcass traits in the pre-weaning and post-weaning periods. Relative to the progeny of N\_N cows, the progenies of crossbred cows tended to have greater birth BW (P=0.058) and significantly greater weaning BW (P<0.001) and pre-weaning ADG (P<0.0001). Nevertheless, the progeny of N\_N cows had greater postweaning P1 and P1 + P2 ADG (P<0.05). Regarding carcass traits, the progeny from crossbred cows tended to have greater B\_T (P=0.08) and had significantly greater R\_T (P<0.05). No differences were observed for post-weaning P2 final BW, post-weaning P2 ADG or R\_A (P>0.05).

Table 5 shows the results of the feedlot performance and ultrasound carcass traits of progenies from different dam and sire breeds, during the finishing period. There were no interaction effects of sire and dam breeds on any evaluated traits (P > 0.05). Dam breed did not affect (P > 0.05) initial or final BW, feed conversion ratio or gross feed efficiency but influenced ADG and DMI ( $P \le 0.05$ ). Progeny of A\_N cows had greater ADG than those of N N cows, whereas C N cows did not differ from either. Progenies from A\_N and C\_N cows had similar DMI values but higher values than those from N\_N cows, both in kg/day and in percentage of BW. Sire breed did not affect (P > 0.05) initial or final BW, DMI, feed conversion ratio or gross feed efficiency but influenced ADG ( $P \le 0.05$ ). Relative to those from Caracu sires, progenies from Braford sires had a greater ADG, whereas those from Charbray sires did not differ from either.

Dam breed did not affect R\_A or R\_T (P > 0.05). Progeny of C\_N cows had lower B\_T (P < 0.05) than those of A\_N cows, but they did not differ from those of N\_N cows (P < 0.05). No sire breed effects were observed for R\_A or B\_T (P > 0.05). However, progeny of Braford sires had

|                        |                          | Dam breed                |                           |         |                          | Sire breed                |                           |         |
|------------------------|--------------------------|--------------------------|---------------------------|---------|--------------------------|---------------------------|---------------------------|---------|
| Variables              | A_N                      | C_N                      | N_N                       | P-value | Braford                  | Charbray                  | Caracu                    | P-value |
| Initial BW (kg)        | 430 (8.0) <sup>2</sup>   | 429 (6.9)                | 424 (8.1)                 | 0.842   | 422 (9.1)                | 432 (8.6)                 | 429 (6.1)                 | 0.571   |
| Final BW (kg)          | 563 (9.5)                | 552 (7.9)                | 546 (9.8)                 | 0.390   | 553 (11.2)               | 561 (10.3)                | 548 (6.8)                 | 0.569   |
| ADG (kg/day)           | 1.47 <sup>a</sup> (0.05) | 1.38 <sup>a</sup> (0.04) | 1.35 <sup>b</sup> (0.05)  | 0.050   | 1.46 <sup>a</sup> (0.06) | 1.43 <sup>ab</sup> (0.05) | 1.31 <sup>b</sup> (0.04)  | 0.045   |
| DMI (kg/day)           | 11.85ª (0.27)            | 11.53ª (0.23)            | 10.93 <sup>b</sup> (0.27) | 0.023   | 11.48 (0.32)             | 11.44 (0.29)              | 11.39 (0.19)              | 0.967   |
| DMI (%BW)              | 2.43 <sup>a</sup> (0.04) | 2.39 <sup>a</sup> (0.04) | 2.27 <sup>b</sup> (0.04)  | 0.002   | 2.39 (0.05)              | 2.34 (0.05)               | 2.35 (0.03)               | 0.712   |
| FCR                    | 8.44 (0.31)              | 8.74 (0.26)              | 8.47 (0.32)               | 0.564   | 8.27 (0.36)              | 8.39 (0.34)               | 8.99 (0.23)               | 0.144   |
| GFE                    | 0.134 (0.01)             | 0.127 (0.01)             | 0.135 (0.01)              | 0.456   | 0.135 (0.01)             | 0.133 (0.01)              | 0.128 (0.01)              | 0.652   |
| R_A (cm <sup>2</sup> ) | 87.7 (2.58)              | 90.5 (2.17)              | 87.8 (2.57)               | 0.445   | 87.6 (2.96)              | 89.8 (2.85)               | 88.6 (1.95)               | 0.805   |
| B_T (mm)               | 9.1ª (0.71)              | 7.4 <sup>b</sup> (0.64)  | 8.2 <sup>ab</sup> (0.73)  | 0.023   | 8.8 (0.79)               | 8.3 (0.75)                | 7.6 (0.59)                | 0.251   |
| R_T (mm)               | 10.7 (0.81)              | 9.4 (0.72)               | 10.5 (0.81)               | 0.135   | 11.3 <sup>a</sup> (0.99) | 9.08 <sup>b</sup> (0.95)  | 10.1 <sup>ab</sup> (0.68) | 0.042   |

**Table 5** Effect of dam and sire breeds on progeny growth performance and ultrasound carcass traits in the finishing period<sup>1</sup>

 $A_N = \frac{1}{2}$  Angus +  $\frac{1}{2}$  Nellore cows,  $C_N = \frac{1}{2}$  Caracu +  $\frac{1}{2}$  Nellore cows;  $N_N$ =Nellore cows; ADG = average daily gain; DMI = dry matter intake; FCR = feed conversion ratio; GFE = gross feed efficiency;  $R_A$  = ultrasound ribeye area;  $B_T$  = ultrasound backfat thickness;  $R_T$  = ultrasound rump fat thickness.

<sup>1</sup> Cattle were generated from mating Nellore, ½ Angus +½ Nellore and ½ Caracu +½ Nellore cows with Braford, Charbray and Caracu sires.

<sup>2</sup> Standard error of mean

<sup>a,b</sup> Values within a row with different superscripts differ significantly at P < 0.05.

| Table   | 6  | Effect  | of  | sire  | breed    | on    | progeny   | feedlo            | ot perf | formance | and |
|---------|----|---------|-----|-------|----------|-------|-----------|-------------------|---------|----------|-----|
| ultraso | un | d carca | ass | trait | s in the | e fin | ishing pe | riod <sup>1</sup> |         |          |     |

|                        | Sire l                 | breed         |         |
|------------------------|------------------------|---------------|---------|
| Variables              | Composite <sup>2</sup> | Caracu        | P-value |
| Initial BW (kg)        | 427 (8.9) <sup>3</sup> | 429 (6.1)     | 0.829   |
| Final BW (kg)          | 557 (10.7)             | 548 (6.8)     | 0.406   |
| ADG (kg/day)           | 1.45 (0.05)            | 1.31 (0.04)   | 0.015   |
| DMI (kg/day)           | 11.46 (0.31)           | 11.39 (0.19)  | 0.824   |
| DMI (%BW)              | 2.37 (0.05)            | 2.35 (0.03)   | 0.879   |
| FCR                    | 8.33 (0.35)            | 8.99 (0.23)   | 0.052   |
| GFE                    | 0.134 (0.008)          | 0.128 (0.005) | 0.367   |
| R_A (cm <sup>2</sup> ) | 88.68 (2.91)           | 88.65 (1.95)  | 0.993   |
| B_T (mm)               | 8.55 (0.77)            | 7.59 (0.59)   | 0.135   |
| R_T (mm)               | 10.20 (0.97)           | 10.13 (0.68)  | 0.945   |

ADG = average daily gain; DMI = dry matter intake; FCR = feed conversion ratio; GFE = gross feed efficiency;  $R_A$  = ultrasound ribeye area;  $B_T$  = ultrasound backfat thickness;  $R_T$  = ultrasound rump fat thickness.

 $^1$  Cattle were generated from mating Nellore,  $^{1\!\!/}_2$  Angus +  $^{1\!\!/}_2$  Nellore and  $^{1\!\!/}_2$ 

Caracu + 1/2 Nellore cows with Braford, Charbray and Caracu sires.

<sup>2</sup> Composite breeds = Braford and Charbray.

<sup>3</sup> Standard error of mean.

greater R\_T than did those from Charbray sires (P < 0.05), and those from Caracu did not differ from either.

Table 6 shows the results from the contrasts approach to compare progenies from composite (Braford and Charbray) and Caracu sires for feedlot performance and ultrasound carcass traits. No differences were observed for initial or final BW, DMI, gross feed efficiency, R\_A, B\_T or R\_T (P > 0.05). Relative to the progeny of Caracu sires, progeny of composite sires had greater ADG (P < 0.05) and tended to have a lower feed conversion ratio (P = 0.052).

## Discussion

Nellore cows and Caracu sires had calves with lower birth weight, a characteristic that can be considered an advantage

due to lower risk of dystocia (Ahlberg *et al.*, 2016). Compared to crossbred cows, Nellore cows may have a uterine environment that limits foetal growth which may partially explain these findings (Ferrel, 1991). This agrees with Calegare *et al.* (2009), Lema *et al.* (2011) and Reggiori *et al.* (2016) who reported that the use of the Nellore breed as a maternal line in a crossbreeding scheme can reduce the birth weight of calves as well as by the study of Perotto *et al.* (1998) who observed lower birth weight in the progeny of Caracu sires compared to those of Charolais sires.

The occurrence of lower weaning weight, due to poor maternal ability or low calf growth potential, has a significant negative effect on cow/calf operations due to lower commercial value of the offspring. When considering sire breeds, no influences were observed on weaning weight, which may suggest that any of the three evaluated breeds may be employed in crossbreeding schemes. On the other hand, when looking at dam breed effects, Nellore cows weaned calves approximately 8% lighter in relation to the crossbred dams, which is confirmed by the results of the contrasts approach. This finding is consistent with regard to differences in birth BW, as well as with regard to pre-weaning growth rate.

Crossbred progenies showed greater weaning BW which may be explained by the superiority of crossbred cows in relation to purebred cows in terms of milk yield as well as by gains obtained through heterosis (Calegare *et al.*, 2007; Rodrigues *et al.*, 2014). On the other hand, the sire breed effects on weaning BW are not supported by the literature, which extensively reports significant differences between British and Continental breeds (Wheeler *et al.*, 2006; Casas *et al.*, 2012).

From the economic point of view, these results may indicate that using Nellore cows may bring lower revenues to producers who depend on the sale of weaned calves when compared to the use of crossbred cows, due to lower weaning BW. However, one should mention that higher resistance to parasites and diseases usually observed in zebu cattle is important to avoid economic losses. Despite the present study not having evaluated health status and treatment of diseases, this may be considered of great concern. In addition, crossbred cows may be heavier than Nellore cows, which results in higher nutritional costs to maintain the herd.

In the context of the present study, the post-weaning period was divided into two phases due to intrinsic pasture-related nutritional differences between them, as previously explained, and confirmed by growth performance results, as the overall performance was 40% lower in P1 compared to P2. The progeny of Nellore cows had ADG values 12% higher than those of crossbred cows during post-weaning phase 1. As post-weaning phase 1 represented the dry season of the year, a better performance in this phase is probably related to the greater degree of adaptation to a lower nutritional level of the animals with a higher proportion of zebu genes (Burrow, 2012).

Regarding sire breed effects, the performance considering the total post-weaning period (P1 + P2) was similar among Caracu, Braford and Charbray. However, during P1 the progeny of Caracu sires had ADG values approximately 14% higher than those of progenies of Braford sires. This difference may be related both to greater adaptability of the Caracu breed and to a greater difficulty of adaptation of the Braford breed, which is derived from a British taurine breed (Hereford). A result that supports this assumption is that Caracu  $\times$  N\_N cattle had a value of ADG 57% higher than that of Braford  $\times$  C\_N.

Although not reflected in significant changes in final feedlot BW, both dam and sire breed effects were observed on feedlot growth rate, where N\_N cows and Caracu sires had progenies with significant lower ADG. This difference in ADG also led to the superiority of the composite breeds for this variable in the contrasts approach. Lower ADG may be partially explained by lower DMI found for the progeny of N\_N cows and by a worse feed conversion ratio of progeny of Caracu sires (contrast approach). From an economic point of view, poor ADG and feed conversion ratios are undesirable, because they are highly correlated with profitability (Cruz, *et al.*, 2010). Therefore, for feedlot-based finishing systems, progeny of composite sires and crossbred cows would be recommended.

Ribeye area was weakly influenced by breed, except for the measure taken at the end of the post-weaning period, when a slight superiority was observed for progeny of Caracu sires compared to progeny of Braford sires. The greater ribeye measurements observed in progeny of Caracu sires relative to those of the Braford sires at the end of the post-weaning period may be related to possible higher heterosis gains obtained from the use of an adapted taurine breed compared to the use of composite breeds in crossbreeding, or to the greater degree of adaptability to tropical pasture-based production system conditions of the Caracu breed relative to Braford, resulting in greater muscular growth. However, such differences were not confirmed at the end of the finishing period following a better nutritional regime compared to the previous production phase.

On the other hand, marked changes in backfat and rump fat were evident due to differences in dam and sire breeds. The higher backfat and rump fat values observed in offspring of N\_N cows relative to offspring of C\_N cows are probably related to the better performance of the former in the total post-weaning period or to a late fat deposition profile of the latter. Moreover, when the contrasts approach was performed, this difference was large enough to establish the superiority of N\_N cows over crossbred cows for rump fat.

No effects of sire breed were observed on B T measures; however, the Braford sire progeny presented higher R T than did those of Charbray. This result may be related to the fact that the Braford breed is partially derived from the British Hereford breed (5/8 Hereford + 3/8 Zebu), which is earlymaturing in regard to the deposition of fat in the carcass relative to the continental taurine breeds such as Charolais. The Charbray breed is partially derived from Charolais (5/8 Charolais + 3/8 Zebu) and is larger in size (frame) and has delayed fat deposition in the carcass relative to that in the Hereford breed. Several studies have shown that continental breeds are larger and heavier than British and Zebu breeds, mature later and have leaner carcasses at the same age (Barton et al., 2006; Albertí et al., 2008; Bidner et al., 2009). Thus, for markets that require greater fat finishing, the present study indicates Braford sires with N N or A N dams as the best alternatives.

In relation to the Nellore cows, higher B\_T values observed at the end of the post-weaning period compared to the C\_N progeny were probably due to the better growth rate in the post-weaning period. The superiority of the A\_N cow progeny when compared to those of C\_N cows in the B\_T measured at the end of the finishing period may be related to the presence of genes from the Angus breed in the former, promoting better fat deposition.

At the end of the finishing period, the progeny of Braford bulls presented higher R\_T in comparison to the Charbray progeny, following the tendency observed at the end of the post-weaning period and confirming the studies reporting the superiority of British breeds relative to continental breeds in fat deposition (Casas and Cundiff, 2006; Prado *et al.*, 2008; Bidner *et al.*, 2009). These results suggest the need to slaughter progeny of Charbray bulls with higher BW, aiming to increase the degree of subcutaneous fat.

It is concluded that in the production system evaluated, the use of crossbred dams promotes increased productivity until weaning, but these benefits are not maintained in the post-weaning and finishing phases. When the objective is to increase the fat finishing rate of the carcasses, the use of  $\frac{1}{2}$  Angus +  $\frac{1}{2}$  Nellore and the use of Nellore cows are considered better options than using  $\frac{1}{2}$  Caracu +  $\frac{1}{2}$  Nellore cows. Regarding the sire breeds evaluated, the use of Braford bulls allows improvements in carcass fat deposition in the finishing period. Considering other carcass variables and performance in the different growing phases, the Favero, Menezes, Torres Jr., Silva, Bonin, Feijó, Altrak, Niwa, Kazama, Mizubuti and Gomes

Braford, Charbray and Caracu breeds produce similar results, though Caracu tends to have inferior weight gain and feed efficiency.

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#### **Declaration of interest**

Authors have no conflict of interest to declare.

## **Ethics committee**

All procedures were approved by Embrapa Beef Cattle Animal Care and Use Committee (#02/2014).

## Software and data repository resources

None of the data were deposited in an official repository.

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