REPRODUCTIVE TRAITS SELECTION IN NELORE BEEF CATTLE

Seleção para características reprodutivas em bovinos de corte da raça Nelore

Heverton Luis Moreira¹, Marcos Eli Buzanskas², Danisio Prado Munari³, Érika Breda Canova¹, Raysildo Barbosa Lôbo⁴, Claudia Cristina Paro de Paz³

ABSTRACT

Genetic breeding programs of beef cattle in Brazil are including new features, mainly related to reproductive efficiency. Thus, it is necessary to study the effectiveness of selection and quantify genetic gain for these traits in herds. This study estimated genetic and phenotypic parameters and genetic trends for reproductive traits used in breeding programs for Nelore beef cattle. The traits studied were the scrotal circumference (SC) at 365 and 450 days of age (SC365 and SC450), age at first calving (AFC) and gestation length, as a cow trait (GLcow) and a calf trait (GLcalf). The (co)variance components were obtained with the Restricted Maximum Likelihood Methodology in a single and double-trait analysis of the animal model. For scrotal circumference (SC365 and SC450), positive and favorable genetic gains were observed. For AFC, GLcow and GLcalf, the trends were favorable for selection, but without significant genetic gain. Selection for large SC may reduce AFC and improve female reproductive efficiency. The selection for reproductive traits (SC365, SC450, AFC and GL) may improve reproductive and productive efficiency of Nelore cattle, if used as a selection criterion.

Index terms: Reproductive efficiency; genetic evaluation; precocity.

INTRODUCTION

New market concepts aim to provide agricultural and livestock products in sustainable production systems. Animal genetic breeding is one of the multidisciplinary tools for this new beef cattle system to meet basic requirements of the sustainable production process. Genetic breeding allows to reduce production cycle, maximizing productive and reproductive potential of the herd, that is, animals with precocious sexual development, growth and carcass finishing. This can improve quality and increase production of Brazilian meat products.

Traits related to reproduction have a great impact on profitability in animal production systems. Thus, selection criteria connected to with sexual precocity and reproductive efficiency have been used widely in genetic breeding programs in Brazil (Lôbo et al., 2010). According to Silva et al. (2011), scrotal circumference (SC) is easily measured and is directly associated with sexual precocity and sperm quantity and quality, which may lead to favorable reproductive trends.
MATERIAL AND METHODS

We used data from animals born between 1998 and 2008, from 22 herds of Nelore Genetic Breeding Program (Nelore Brazil), coordinated by the National Breeders and Researchers Association (ANCP). All animals in the program undergo zootechnical monitoring of reproductive and productive traits and are kept in an extensive or semi-extensive feeding regimen. The reproductive management consisted of artificial insemination of females during the mating period, when they were kept in corrals with clean-up bulls.

In males, SC was measured in centimeters (cm), corrected to 365 days of age (SC365) and 450 days of age (SC450). In females, AFC was measured in months, obtained from the difference between the birth date and date of first calving; GL was measured in days, obtained from the difference between insemination date and calving date, which was obtained for cow trait (GLcow) and calf trait (GLcalf).

Contemporary groups (CG) were defined based on the analysis performed in the least square method, using the GLM procedure of the SAS software (SAS, 2002-2003), to define the fixed effects in the mixed models. Farm data (Nfa), year of birth (YoB), season of birth (SoB), defined as 1 = dry (April to September) and 2 = rainy (October to March), management batch (Batch 120, 365 and 450 days), Sex (Sx) and dam age at calving (Dac) significantly affected (P<0.005) the studied traits. CG were defined as SC365: Nfa + YoB + SoB + Batch120 + Batch365; SC450: Nfa + YoB + SoB + Batch120 + Batch365 + Batch450; AFC: YoB + SoB + Batch120 + Batch365 + Batch450; GLcow: Sx + Nfa + Dac; GLcalf: Sx + Nfa + YoB + SoB + Dac.

Normality of residuals was carried out in the Cramer-von Mises test, however, normality was not observed for the studied data. Standardized residuals were used to evaluate discrepant data. Observations where residuals were higher than 3.5 standard deviations and lower than -3.5 standard deviations were excluded.

After ensuring data file consistency, descriptive statistical estimates were produced for the following traits: SC365 (cm), SC450 (cm), AFC (months), GLcow (days) and GLcalf (days) (Table 1).

The analyses of variance component estimation were performed with REML using the animal model. The fixed effect (CG), random effect (direct additive) and residual error effect were considered for all traits. The covariable of dam age at calving had significant effect (P<0.005) on the studied traits.
Reproductive traits selection...

linear and quadratic (P<0.05) values for SC365, SC450, GLcow and GLcalf. For SC365, SC450 and AFC, the model included only random and direct additive genetic effects. For GLcow, the model included direct additive genetic effects and permanent environment. For GLcalf, the model included random direct additive and maternal genetic effects.

The estimates for (co) variance components, genetic parameters and breeding values for the traits studied were obtained in REML, in a multi-trait animal model, using WOMBAT computer software (Meyer, 2007).

The annual genetic gains of SC365, SC450, AFC, GLcow and GLcalf were estimated in linear regression of breeding value (BV) based on year of birth. The significance of Geometric Growth Rate - GGR was evaluated using the “t” test. The GGR values indicate the mean linear change in the breeding value (%) for each variable.

RESULTS AND DISCUSSION

The estimates of variance components and direct and maternal heritability for the traits studied (SC365, SC450, AFC, GLcow and GLcalf) are shown in Table 2.

Heritability estimations obtained for SC measured at 365 and 450 days of age were 0.41 and 0.44, respectively, similar to estimations described in the literature that, ranged from 0.42 to 0.53 (Pereira; Eler; Ferraz, 2002; Garnero et al., 2002; Dias; El Faro; Albuquerque, 2003; Gianlorenço et al., 2003; Yokoo et al., 2007). These values were very similar, indicating that the used of the same selection intensity at both ages favors the response for the selection of SC365 because genetic gains of this trait are associated to economic returns, with the choice of sexually precocious bulls fit for the breeding season.

Heritability for AFC was 0.20, showing that this value can be related to the analysis method that considered only females with their first birth on the farm or other factors such as nutrition and mating criteria used for steers exposure as cited by Pereira, Eler and Ferraz (2002) and Dias, El Faro and Albuquerque (2004). Therefore, AFC genetic variability can be obtained from direct selection, however, the methods cited above should be improved in order to explore the trait variability, which could bring direct economic efficiency, profitability and competitiveness of Brazilian cattle (Boligon; Rorato; Albuquerque, 2008).

Table 1: Number of animals, number of contemporary groups (CG), mean, standard deviation (SD), minimum (Min) and maximum (Max) for the analyzed traits.

<table>
<thead>
<tr>
<th>Trait*</th>
<th>Animals</th>
<th>CG</th>
<th>Mean</th>
<th>SD</th>
<th>Min</th>
<th>Max</th>
</tr>
</thead>
<tbody>
<tr>
<td>SC365 (cm)</td>
<td>18747</td>
<td>837</td>
<td>20.43</td>
<td>2.25</td>
<td>12.30</td>
<td>30.00</td>
</tr>
<tr>
<td>SC450 (cm)</td>
<td>18430</td>
<td>924</td>
<td>23.64</td>
<td>3.06</td>
<td>12.80</td>
<td>35.90</td>
</tr>
<tr>
<td>AFC (months)</td>
<td>13329</td>
<td>875</td>
<td>35.16</td>
<td>4.98</td>
<td>21.00</td>
<td>49.00</td>
</tr>
<tr>
<td>GLcow (days)</td>
<td>22382</td>
<td>2335</td>
<td>296.49</td>
<td>5.85</td>
<td>277.00</td>
<td>314.00</td>
</tr>
<tr>
<td>GLcalf (days)</td>
<td>12283</td>
<td>2427</td>
<td>296.66</td>
<td>5.75</td>
<td>277.00</td>
<td>314.00</td>
</tr>
</tbody>
</table>

* Scrotal circumference at 365 (SC365) and 450 (SC450) days of age, age at first calving (AFC), gestation length as a cow trait (GLcow) and gestation length as a calf trait (GLcalf).

Table 2: Additive (σ²a), residual (σ²e) and phenotypic (σ²p) genetic variance components, maternal additive variance (σ²am), maternal variance (σ²m), permanent environment (σ²ep), direct heritability with standard error (h²d±SE) and maternal heritability with standard error (h²m±SE), in a single-trait analysis.

<table>
<thead>
<tr>
<th>Trait*</th>
<th>σ²a</th>
<th>σ²e</th>
<th>σ²am</th>
<th>σ²m</th>
<th>σ²ep</th>
<th>h²d±SE</th>
<th>h²m±SE</th>
</tr>
</thead>
<tbody>
<tr>
<td>SC365</td>
<td>120.37</td>
<td>171.62</td>
<td>----</td>
<td>----</td>
<td>----</td>
<td>0.41±0.02</td>
<td>----</td>
</tr>
<tr>
<td>SC450</td>
<td>226.60</td>
<td>286.14</td>
<td>----</td>
<td>----</td>
<td>----</td>
<td>0.44±0.02</td>
<td>----</td>
</tr>
<tr>
<td>AFC</td>
<td>3.12</td>
<td>12.50</td>
<td>----</td>
<td>----</td>
<td>----</td>
<td>0.20±0.02</td>
<td>----</td>
</tr>
<tr>
<td>GLcow</td>
<td>6.14</td>
<td>23.21</td>
<td>----</td>
<td>----</td>
<td>1.40</td>
<td>0.20±0.01</td>
<td>----</td>
</tr>
<tr>
<td>GLcalf</td>
<td>14.01</td>
<td>12.62</td>
<td>-0.62</td>
<td>2.67</td>
<td>----</td>
<td>0.49±0.05</td>
<td>0.09±0.02</td>
</tr>
</tbody>
</table>

* Scrotal circumference at 365 (SC365) and 450 (SC450) days of age, age at first calving (AFC), gestation length as a cow trait (GLcow) and gestation length as a calf trait (GLcalf).
The GL evaluated in this study aimed to show if it should be explored in the analysis as genetically influence of the cow (GLcow) or calf (GLcalf), where the coefficient of heritability was larger for GLcalf (0.49 ± 0.05) compared to GLcow (0.20 ± 0.01), which corroborates the study of Mucari et al. (2011) in Canchim. Thus, the GL analyzed as a calf trait could result in greater genetic gain by direct selection, reducing the period, increasing the success of pregnant females in the breeding season (Pereira; Eler; Ferraz, 2002), thus, justifying its inclusion into genetic breeding programs.

The genetic and phenotypic correlations between the traits studied are presented in Table 3. The genetic correlation between SC365 and SC450 (0.96) indicates that most genes responsible for SC at 365 days of age also influence at 450 days. This result is in agreement with Gressler et al. (2000) who found a genetic correlation between SC365 and SC450 of 0.99 and 0.72 phenotypic. Thus, in the selection of bulls with larger SC365, the correlated response at SC450 is the same.

Genetic correlation estimates between SC365 and SC450 were then associated to AFC (Table 3), and the results obtained were -0.40 at SC365 from AFC and -0.37 at SC450 from AFC, indicating that bull selection based on larger SC at both ages reduces AFC of their female offspring. The literature reported negative results of associations SC and AFC in Nelore ranging from -0.22 to -1.00 (Gressler et al., 2000; Pereira; Eler; Ferraz, 2000; Pereira; Eler; Ferraz, 2002; DIAS et al., 2003). This shows that the indirect gains of selection to promote SC at first calving, which is economically viable for beef cattle production systems, because measuring SC is somewhat easily for a low cost.

Genetic correlations between SC365/SC450 days and GLcalf were favorable and negative -0.21 and -0.19 respectively, suggesting that selection for SC could reduce GLcalf, but not in GLcow, where the results were virtually null. AFC also showed favorable genetic associations with GLcalf (0.28) and GLcow (0.19), where gains from selection for reduced AFC also promoted positive impacts on GL. Therefore, the GL as a calf trait had the highest favorable genetic associations with the PE and IPP, that is, selection of bulls with larger SC and females with early AFC could promote indirect reduction of gestational period of the mother, bringing all benefits to improve production systems.

Estimates of genetic trends allow to evaluate how the selection process changes over the years, considering the mean breeding value. Thus, these estimates test the efficacy of the proposed selection indexes for the traits selected. In this study, the selection applied to the males to obtain animals with greater SC at 365 (Figure 1A) and 450 (Figure 1B) days of age, provided genetic gains of 0.025cm/year and 0.034cm/year (P < 0.0001), respectively.

This trend is very clear regarding genetic gain for SC. An increase of 0.12% (SC365) and 0.14% (SC450) per year was observed, in relation to the expected mean for Nelore cattle during the period evaluated. This result was lower than the findings of Cyrillo et al., (2001), who compared a herd under constant selection with a control herd. The animals in the selection herd had an increase of 0.31cm/year, whereas the animals in the control herd showed a decrease of 0.21cm/year for the trait. Therefore, direct selection for bulls with larger SC promoted positive changes, in the studied period, to breeding values of Nelore, which is desirable for programs that uses selections as a criterion, considering their genetic associations favorable with other economically important traits for production systems mentioned in this study.

<table>
<thead>
<tr>
<th>Trait*</th>
<th>SC365</th>
<th>SC450</th>
<th>AFC</th>
<th>GLcow</th>
<th>GLcalf</th>
</tr>
</thead>
<tbody>
<tr>
<td>SC365 (cm)</td>
<td>----</td>
<td>0.96 ± 0.01</td>
<td>-0.40 ± 0.01</td>
<td>0.02 ± 0.01</td>
<td>-0.21 ± 0.01</td>
</tr>
<tr>
<td>SC450 (cm)</td>
<td>0.85 ± 0.01</td>
<td>----</td>
<td>-0.37 ± 0.01</td>
<td>-0.01 ± 0.01</td>
<td>-0.19 ± 0.01</td>
</tr>
<tr>
<td>AFC (months)</td>
<td>0.16 ± 0.01</td>
<td>-0.04 ± 0.01</td>
<td>----</td>
<td>0.19 ± 0.01</td>
<td>0.28 ± 0.01</td>
</tr>
<tr>
<td>GLcow (days)</td>
<td>0.02 ± 0.01</td>
<td>0.01 ± 0.01</td>
<td>0.04± 0.01</td>
<td>----</td>
<td>----</td>
</tr>
<tr>
<td>GLcalf (days)</td>
<td>0.05 ± 0.01</td>
<td>0.01 ± 0.01</td>
<td>0.05± 0.01</td>
<td>----</td>
<td>----</td>
</tr>
</tbody>
</table>

* Scrotal circumference at 365 (SC365) and 450 (SC450) days of age, age at first calving (AFC), gestation length as a cow trait (GLcow) and gestation length as a calf trait (GLcalf).
Genetic gain for AFC (Figure 2) was -0.01 months/year ($P < 0.0001$), and although practically null, it still represents a yearly decrease of 0.02%, showing indirect gains for the trait through selection. The result of low yearly genetic gain for AFC may have reflected on management methods used on farms and genetic evaluation models, and these reductions do not express the potential that breeders expect for females.

The genetic trend for AFC in female Gir cattle showed an increase of 0.008 months/year (Balieiro et al., 1999). Júnior et al., (2010), also studied Gir cattle and found favorable genetic trends, with a yearly gain of -0.018 months. Laureano et al., (2011) studied beef cattle and found a reduction of 1.5 days/year for female Nelore cattle that were challenged precociously. Thus, studies show that estimates for genetic trends have been favorable regarding the genetic potential of sexual precocity, although without significant genetic progress for age at first calving.

Genetic trend estimates for GL considered cow and calf as traits are presented in Figure 3. The genetic change in GL, considering cow responsible for the period, was -0.02 days/year, equal to a mean genetic gain of -0.006% per year. When the gestation period considered a calf trait, the genetic trend was -0.008 days/year, equal to -0.002% per year.

Figure 1: Mean predicted breeding value variation for: (A) scrotal circumference at 365 days of age (SC365), and (B) scrotal circumference at 450 days of age (SC450). The regression coefficients were significantly different ($P < 0.001$) from zero using the t test.

Figure 2: Mean predicted breeding value variation for age at first calving (AFC). The regression coefficients were significantly different ($P < 0.001$) from zero using the t test.
These practically null gains were expected for AFC and GL (cow and calf) traits, since they have become part of the genetic selection index in the Nelore Brazil Program since 2008. Over the period analyzed in the current study, these traits were not part of direct or indirect selection process, which explains the non-significant genetic gains for GL and AFC. From 2008, these traits were included in the selection index, and therefore the genetic gains for GL and AFC in the coming years are different. This suggests that selection for SC (365 and 450), AFC and GL are important in Nelore cattle genetic breeding programs, as these traits may influence economic and socio-environmental issues of the Brazilian beef cattle production and maximize the objectives of selection for sexual precocity in animal breeding systems.

CONCLUSIONS

Direct selection for SC365 and SC450 traits is expected to provide greater genetic gains than the reproductive traits evaluated in females (AFC and GLcow). The resultant gains are favorable through the positive genetic trend rates.

For AFC, greatly influenced by the environment, environmental improvements related to reproductive management will allow females to be better evaluated and consequently selected, possibly increasing their reproductive rates.

Direct selection of GL as a calf trait may bring greater genetic gain than when selected as a cow trait.

The inclusion AFC in the selection of objectives associated with information at SC365 as a selection criterion can promote greater economic response, due to its economic impact on beef cattle production systems.

REFERENCES


Reproductive traits selection...


