

Fixed environmental effects and connectedness of the genetic evaluation of the Limousin breed in Uruguay

Editor

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Efectos ambientales fijos y conectividad de la evaluación genética de la raza Limousin en Uruguay

Efeitos ambientais fixos e conectividade da avaliação genética da raça Limousin no Uruguai

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Abstract

Genetic evaluations are an efficient tool to select sires, and for them it is necessary to have identification of environmental effects, genetic parameters, and acceptable degree of connectedness between herds. Evaluation of Limousin breed genetics has been developing since 2012. The objectives were to determine the most relevant fixed environmental effects of the characteristics in the genetic evaluation and to estimate the degree of connectedness between herds. The productive and genealogical records were extracted from the Limousin genetic evaluation database of the National Institute of Agricultural Research. Objective measurements were available for the characteristics birth weight (916 records), weaning (818), 18 months (438), ribeye area (435), and 12th-13th rib fat thickness (435) for 2012, 2013, 2014, 2015 and 2016 corresponding to four herds. The results showed that the fixed environmental effects considered presented a significant value (p<0.05), and that herds were connected to each other with average values of prediction error variance (PEVD), and connectedness rating (CR) of 113.9 and 7.9, respectively. Concluding that the most relevant fixed environmental effects were year, sex, herd, age at weaning and age at 18 months, and that herds were connected with a low connectedness degree.

Keywords: Limousin, environmental effects, genetic connectedness, Uruguay

Resumen

Desde 2012 la raza Limousin se encuentra en proceso de desarrollo de la evaluación genética poblacional. Dichas evaluaciones genéticas son una herramienta eficiente para seleccionar reproductores, y para las mismas es necesario contar con identificación de efectos ambientales, parámetros genéticos y niveles aceptables de conexión entre cabañas. Los objetivos de este trabajo fueron la determinación de los efectos ambientales fijos más relevantes de las características en la evaluación genética, y estimar el grado de conectividad entre cabañas. Los registros productivos y genealógicos fueron extraídos de la base de datos de la evaluación genética de la raza Limousin por medio del Instituto Nacional de Investigación Agropecuaria. Se dispuso de mediciones objetivas para las características peso al nacimiento (916 datos), destete (818 datos), los 18 meses (438 datos), área del ojo de bife (435 datos) y espesor de grasa subcutánea (435 datos) para 2012, 2013, 2014, 2015 y 2016, correspondientes a cuatro cabañas. Los resultados indican que los efectos ambientales considerados presentaron significancia estadística (p<0,05), y las cabañas se encontraban conectadas entre sí con valores promedio de varianza del error de predicción (PEVD) y ranking de conectividad (CR) igual a 113,9 y 7,9, respectivamente. Se concluye que los efectos ambientales fijos más relevantes fueron año, sexo, cabaña, edad al destete y edad a los 18 meses, y que las cabañas se encontraban conectadas con valores bajos de conectividad.

Palabras clave: Limousin, efectos ambientales, conectividad genética, Uruguay

Resumo

A raça Limousin desde 2012 está em processo de desenvolvimento da avaliação genética de sua população. Estas avaliações genéticas são uma ferramenta eficiente para selecionar os criadores, e para o mesmo, é necessário identificar os efeitos ambientais, parâmetros genéticos e níveis aceitáveis de conexão entre os rebanhos. Os objetivos deste trabalho foram determinar os efeitos ambientais fixos mais relevantes das características na avaliação genética e estimar o grau de conectividade entre os rebanhos. Os registros produtivos e genealógicos foram extraídos do banco de dados de avaliação genética da raça Limousin pelo Instituto Nacional de Pesquisas Agrícolas. Medidas objetivas estavam disponíveis para as características de peso ao



nascer (916 dados), desmame (818 dias), 18 meses (438 dados), área do olho de bife (435 dados) e espessura da gordura subcutânea (435 dados) para 2012, 2013, 2014, 2015 e 2016 para quatro rebanhos. Os resultados indicam que os efeitos ambientais considerados foram estatisticamente significativos (p<0,05), e as cabanas foram conectadas entre si com valores médios de variância do erro preditivo (PED) e ranking de conectividade (CR) iguais a 113,9 e 7,9, respectivamente. Concluímos que os efeitos ambientais fixos mais relevantes foram ano, sexo, rebanho, idade de desmame e idade de 18 meses, e que os rebanhos estavam conectados com baixos valores de conectividade.

Palavras-chave: Limousin, efeitos ambientais, conectividade genética, Uruguai

1. Introduction

Genetic evaluations are tools for the objective selection of sires that allow achieving genetic progress in production animals. In Uruguay, the number of herds and animals participating in genetic evaluations has grown steadily since the first published ones in 1993. Uruguay's Limousin Breeder Society, the Veterinary College, and the National Institute of Agricultural Research (SCLU, FVET and INIA, by their Spanish acronyms) have carried out the genetic evaluation of the breed in Uruguay since 2012. The objective is to achieve a population genetic evaluation, for which it is necessary to have the identification of the environmental factors that affect the evaluated characteristics, the genetic parameters estimated for the population, and the acceptable connectedness levels between the participating herds.

Genetic evaluations predict the genetic merit of an animal, based on its phenotype and/or that of its relatives. Therefore, it must be corrected by known factors that influence its expression, such as nongenetic or environmental effects⁽¹⁾. According to Cardellino and Rovira⁽²⁾, the environment represents those external conditions that affect the reproduction, production, and composition of the cattle carcass, being its influence dependent on the analyzed characteristic. The adequate definition of contemporary groups (CG) is key to correct by environmental effects in a genetic evaluation⁽³⁾, and they were defined by Cundiff and others⁽¹⁾ as the groups of animals of the same breed, sex and who have similar ages and management.

It is necessary to confirm the connectedness between herds, because it allows comparing animals

in different productive environments and making precise comparisons of the breeding values of the candidates for selection(4), thus increasing the intensity of selection and therefore the genetic progress. The easiest way to achieve connectedness within CG is through the use of reference animals, which have progeny with productive records in more than one herd and/or year. The offspring of a sire will share the same genetic information, regardless of the environment they are in(5). There are two main methods for estimating connectedness, firstly, qualitative methods that identify only connected and disconnected groups, and secondly, quantitative methods that determine the degree of connectivity between herds. According to Magaña and others(6), the latter are mostly based on the accuracy of differences in breeding values between two individuals and is expressed as the prediction error variance of the differences (PEVD). According to Kennedy and Trus(7), the most appropriate measure of connectivity is the average PEVD of breeding values of animals from different herds. As there is a large number of animals in genetic evaluations, this measure must consider all pairs of individuals between herds, which makes it too complex for use. Mathur and others(8) proposed an alternative measure of connectivity, which implies the measurement of the connectivity ranking (CR), defined as the correlation between the estimates of the herd effects or CG.

The objectives of this study are to determine the most relevant fixed environmental effects, and to estimate the connectedness rating between herds for the future population genetic evaluation of the Limousin breed in Uruguay.



2. Material and Methods

2.1 Data

The productive and genealogical records used come from the Limousin genetic evaluation database of the SCLU-FVET-INIA agreement.

The analyzed characteristics were weight at birth (WB), weight at weaning (WW), weight at 18 months (W18), ribeye area (REA) and rib fat thickness (RFT) of bulls and heifers born between 2012 and 2016, belonging to four herds.

The data edition consisted of eliminating animals without date of birth and animals that received management based on natural field + ration, or pasture + ration. In turn, outlier records (less than 0.5% of the data) for W18 (>510 kg) and rib fat thickness (>8 mm) were deleted. Animals with weaning ages between 120 and 300 days were included for the characteristic ww (limits established for the national genetic evaluation), and for W18, animals of 457 to 640 days of age were included (18 months ± 3 months). The edition was carried out using the statistical program SAS⁽⁹⁾.

2.2 Statistical analysis

Descriptive statistics of environmental effects were performed using the MEANS procedure of the statistical program SAS⁽⁹⁾. The effects of sex, herd, year of birth, management code, mother's age at parity, age at weaning, and age at 18 months were analyzed.

The study of the fixed effects on growth and carcass quality was carried out with the following model:

$$y_{ijklm} = S_i + C_j + A_k + M_l + P_m + e_{ijklm}$$

where:

 y_{ijklm} = Characteristic for the m-th animal (WB, WW, W18, REA, RFT)

S_i= Sex: i = M. H

 C_i = Herd; j = 1...4

 A_k =Year of birth; k = 2012...2016.

 M_l = Management code; l = 1 ... 3 (1 = only natural field, 2 = natural field + pasture, 3 = only pasture)

 P_m = Mother's age at parity; m = 4 (2 to 4 years), 5, 6, 7, 8 years, 9 (9 to 10 years), x (mother's age unknown)

eijklm = Residual Random Error

Regarding ww, the calf's age at weaning was included in the model as a covariate, and for w18, REA and RFT the covariate age at 18 months. Possible interactions between effects were not considered due to the small number of records.

The analysis of fixed environmental effects included in the model was performed using the GLM procedure of the statistical program SAS⁽⁹⁾.

2.3 Connectedness

The study of connectedness between herds and years considered the same data as the analysis of environmental effects, and had information about the reference parents used.

The program CONEXF90⁽¹⁰⁾ was used to determine PEVD and CR, considering weaning weight data grouped by year and CG, and assuming a heritability value of 0.30 for this characteristic according to the studies presented by Gregory and others⁽¹¹⁾, and Phocas and Laloë⁽¹²⁾.

3. Results and discussion

3.1 Environmental effects

The results of the descriptive statistics (Table 1) presented differences between males and females for all the characteristics, resulting in significant differences (p <0.05) for the characteristics, except for RFT. The results of the model for growth and carcass quality are presented in Table 2.



Table 1. Descriptive statistics of the sex effect for the characteristics: birth weight, weaning weight, weight at 18 months, ribeye area, and rib fat thickness.

		Males					Females					
Characteristic	Average	SD	Min.	Max.	n	Average	SD	Min.	Max.	n		
Birth Weight (kg)	37.4	3.9	20.0	52.0	456.0	35.7	3.7	21.0	49.0	460.0		
Weaning Weight (kg)	173.7	37.9	75.0	339.0	390.0	166.3	31.9	85.0	290.0	428.0		
Weight at 18 months (kg)	327.9	50.3	120.0	437.0	217.0	268.9	44.5	104.0	393.0	221.0		
Ribeye Area (cm2)	56.7	9.8	26.4	89.7	215.0	44.2	7.6	25.0	91.2	220.0		
Rib Fat Thickness (mm)	1.8	0.3	1.0	2.9	215.0	1.8	0.3	1.0	2.9	220.0		

SD = Standard deviation, n = number of records

Table 2. Variance analysis for birth weight, weaning weight, weight at 18 months, ribeye area (rea), and rib fat thickness (rft) of Limousin animals.

	R ²	Sum of squares (type III)											
Characteristic		Herd		Year		Sex		Mother's age		Management code		Weaning age	
Birth weight	0.13	126.7	*	288.1	**	704.2	**	349.9	**	1.1	ns		
Weaning weight	0.31	57736.6	**	23173.7	**	9725.6	**	26236.7	**	63.5	ns	185.991.6	**
Weight at 18 months	0.51	98740.0	**	48151.6	**	172047.1	**	28957.9	*	3878.9	ns		
REA	0.51	3312.5	**	1641.2	**	6311.7	**	732.6	ns	383.7	*		
RFT	0.09	0.6	ns	1.6	**	0.1	ns	0.3	ns	0.5	*		

ns= not significative. *=P <0.05; **P <0.01.

The sex effect presented a significative value for the characteristics: WB, WW, W18 and REA, which coincides with the results of Simcic and others⁽¹³⁾, that worked with 167 Limousin animals, and Koch and Clark⁽¹⁴⁾, who with 5952 Hereford breed animals obtained that males were 1.2 kg and 11.8 kg heavier than females at birth and at weaning, respectively. Goldberg and Ravagnolo⁽¹⁵⁾, with Angus animals in Uruguay, found that the effects herd, year, and month of measurement of the characteristic were significant for the characteristics WB, WW, and W18. In this study with the Limousin breed, the herd and year effects significantly influenced all the characteristics, except for the herd effect for RFT.

For their part, the Limousin calves born from 6 and 9-year-old mothers in this study were 1.9 kg (p<0.05) and 2.7 kg (p<0.01) heavier at birth than

those from 2 to 4-year-olds, respectively. Meanwhile, 5, 7, and 8-year-old mothers had heavier calves at birth, but the differences were not significant with respect to 2 to 4-year-olds. Krupa and others⁽¹⁶⁾ reported similar results with various breeds, where 5 to 7-year-old mothers delivered the heaviest calves at birth. Koch and Clark⁽¹⁴⁾ obtained higher WB and WW in calves born to 3 to 6-year-old mothers. The inclusion in this work of a category of mothers where mothers with different ages were grouped because they did not have a date of birth could explain the differences with these authors.

The management effect was not significant to growth characteristics (WB, WW, and W18), which was striking given the differences in animal performance normally observed nationally when natural field, natural field and pasture, and only pasture



managements are applied in the conditions of our country.

Weaning age covariate was significant for ww, where for each day animals presented 0.48 kg more at weaning (p <0.001), while covariate age at 18 months did not present statistical significance for w18, REA and RFT.

3.2 Connectedness

From the information of the sires used by the herds and the effective offspring obtained in each herd, they were directly and/or indirectly connected only in the years 2013, 2014 and 2015. For 2012 there were only two herds, which were connected by a single parent. The number of sires used in common for the remaining years was three to four, with

an average of six offspring per herd. In 2016, three herds were connected through two parents in common.

The average, minimum and maximum values of PEVD of a herd with the others were 113.9, 27.8, and 354.3, respectively. Lower values of PEVD would indicate a greater connectivity ranking between herds. PEVD values from 2012 to 2016 tend to the reduction of these towards this last year (Table 3).

Table 3 shows PEVD values and the connectedness rating (CR) between herds. The correlation between CR and PEVD is -0.45⁽¹⁷⁾. The average, minimum and maximum values of CR of a herd with the other herds were 7.9, 0.03 and 58.6, respectively.

Table 3. Number of animals, minimum, maximum and herd which presents the highest connectivity ranking (cr), and prediction error variance average (pevd) per herd and year.

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Year	Herd	Number of	PEVD	CR	(%)	Herd with greater connectedness	
i cai	Tibiu	animals	Average	Minimum	Maximum		
2012	С	34	112.7	0.6	14.8	Α	
	Α	20	145.3	0.4	14.8	С	
12	D	33	121.9	0.3	7.0	С	
	В	15	263.3	0.3	0.6	С	
	В	5	111.9	4.6	11.9	С	
20	С	65	99.4	3.3	11.9	В	
2013	D	67	195.2	1.1	5.6	В	
	А	47	137.2	1.1	4.6	В	
	А	45	86.1	1.9	9.6	С	
2014	С	70	93.7	2.2	9.6	Α	
114	D	96	122.2	1.1	3.3	С	
	В	20	77.1	1.1	2.2	С	
2015	В	33	99.4	1.5	9.1	D	
	D	66	171.2	0.6	9.1	В	
	С	21	107.0	1.2	5.3	В	
	А	39	107.0	0.6	1.5	В	
2016	С	8	95.0	2.2	9.8	Α	
	А	6	74.4	1.8	9.8	С	
	В	12	81.0	3.3	6.0	С	
	D	74	66.1	1.8	3.7	В	



There are few references to CR values in the literature. Mathur and others⁽⁸⁾ worked with rear fat thickness in swines, and propose a connectivity ranking greater than 3%, as an acceptable CR value. Considering that value as a reference, the A, B, and C Limousin herds for 2016 have an acceptable degree of connectedness between them.

For their part, Roso and others⁽¹⁷⁾ studied different methods of measuring connectivity in bulls from a Canadian testing center They obtained an average value of PEVD of 1726 between a group of animals and the rest, and a CR of 1.2%. Regarding the work carried out with the Limousin breed in Uruguay, the results are different for both measures, since average values of 113.9 and 7.9% were obtained for PEVD and CR, respectively.

The results could be explained by the small size of the contemporary groups considered in the study since, as mentioned by Mathur and others⁽⁸⁾, it is recommended that the GC have at least ten animals and three different parents to obtain adequate connectedness values. Roso and others⁽¹⁷⁾ state that a group evaluated with few parents presents a greater variation in the PEVD and CR, which indicates a greater variation in the degree of connectedness between groups. It is important to note that the smaller the number of animals per group, the PEVD increases and the CR decreases.

Based on what was presented, the herds participating in the genetic evaluations of the Limousin breed for the evaluated years were directly and/or indirectly connected, but with low connectivity values (less than 3% of CR proposed by Mathur and others⁽⁸⁾) that did not yet warrant a population genetic evaluation.

4. Conclusions

The fixed environmental effects of year, sex, herd, age at weaning, and age at 18 months are relevant and should be included in studies associated with the genetic evaluation of the Limousin breed. In order to consolidate the population genetic evaluation of the breed in Uruguay, it is necessary to improve the connectedness rating between herds by using reference parents, and to present contempo-

rary group sizes greater than or equal to 10 animals.

Author contribution statement

G.M.B. conceived and designed the analysis, performed the analysis, and wrote the paper; R.L. collected the data and contributed to data analysis; F.M. collected the data and contributed to data analysis; M.L. collected the data and contributed to data analysis; O.R. collected the data, conceived and designed the analysis, and contributed to data analysis.

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