

Original Research Paper

Evaluation of Relationships Between Some Growth Traits Measured at Birth and Weaning in South African Non-Descript Goat Kids using Canonical Correlation Analysis

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Abstract: The study was conducted to assess the association among 6 different growth traits viz; X set- Birth Weight (BW), Heart Girth (HG), Rump Height (RH), Sternum Height (SH), Body Length (BL), Withers Height (WH), assessed at birth and viz; Y Set- Heart Girth (HG_3), Body Length (BL_3), Rump Height (RH_3), Weaning Weight (WW), Withers Height (WH_3), Sternum Height (SH_3) at weaning Measured in fifty (n = 50) non-descript goat kids. Pearson's correlation and canonical correlation analysis were employed, outcome of Pearson's highlighted that BW had a positive highly statistical correlation with WW ($r = 0.85^{**}$), WH_3 ($r = 0.61^{**}$) and SH_3 ($r = 0.83^{**}$) at $P < 0.01$. Results of canonical correlation showed an estimated canonical correlation coefficient ($r = 0.881$) that was significant among the first pair of canonical variables ($P < 0.01$). The results indicated that WH and HG had a greater contribution to the explanatory capacity of canonical correlation estimation at birth while BL_3 and RH_3 contributed mostly at weaning of non-descript goats. The findings of the study suggest that the heart girth and withers height might be used at birth as criteria for genetic improvement in non-descript goat kid's selection.

Keywords: Birth Weight, Canonical Weights, Canonical Variable, Canonical Correlation Coefficient

Introduction

South African non-descript goats are important to the communal farmers as they provide income and demand less capital for treatments and feed-supplement as they are hardy, resistant to diseases, good forager, well adapted to the harsh environment and fertile from a young age (Tyasi *et al.*, 2020; Norris *et al.*, 2015). Economic important traits that can be used in selection for improvement of the herd include bodyweight (Hagos, 2016; Iqbal *et al.*, 2013; Tsegaye *et al.*, 2013). Determining the association between two or more traits at birth and weaning is important in improving selection (Sahin *et al.*, 2011). Many researchers prefer the simple correlation to determine the association and degree of association between body measurement traits (Berhe, 2017; Patbandha *et al.*, 2018). However, the interrelationship among one or more variables makes it difficult to pinpoint the significance of the correlation (Tahtali *et al.*, 2012). Çankaya and Kaylaap, (2007) define canonical correlation as the technique used to

describe the relation between two sets of variables by calculating a linear combination that is having maximum correlation. The canonical correlation has been used to describe the relationship among birth weight and weaning weight in German sheep breeds (Çankaya and Kaylaap, 2007), reproduction and production traits of meat-type quails (Ribeiro *et al.*, 2016), milk and wool yield in Akkaraman sheep in previous livestock studies (Karadavut, 2020). However, to our knowledge, there is no study estimating the relationship using the canonical correlation technique in Non-descript goats. Hence, the objectives of the paper were to (1) investigate the interrelationship between the six different body measurement traits at birth and weaning; (2) to explore which traits might be used at birth as selection criteria to improve the live body weight of non-descript goats at birth and weaning. This study will assist small stock farmers to identify interrelation among growth traits and making a good decision when selecting traits to improve the live body weight at birth and weaning.

Material and Methods

The research was accomplished in a research area as outlined by Alabi *et al.* (2012). The experimental animals were a total of fifty (50) South African non-descript goat kids. A weighing scale was used to measure Body Weight (BW) while growth traits, namely Body Length (BL), Sternum Height (SH), Heart Girth (HG), withers height (WH) and Rump Height (RH) were measured at birth and weaning using a tape and a wooden ruler calibrated in centimetres. The measuring of the entire growth traits was carried out as described by Norris *et al.*, 2015. Birth weight was measured after 24 hours and weaning weight was measured at three months of age. All the goat kids were weighed 24 hours after birth and ear-tagged. The goat kids were kept with their dam in the kraal with other goats. The kids were left in the kraal with does overnight; the only time the kids were separated from the does was when the does were released for grazing.

Canonical Correlation Analysis (CCA)

Canonical correlation technique was performed as described by Tahtali *et al.* (2012). Briefly, body measurement traits at birth were set as X-Set while body measurement traits at weaning were set as Y-Set. Interpretation of the canonical correlation was performed following the procedure of Sahin *et al.* (2011).

The null and alternative hypotheses for assessing the statistical significance of the canonical correlation coefficients are:

$$H_0: \rho_1 = \rho_2 = \dots = \rho_r = 0$$

$$H_1: \rho_i \neq 0 \text{ I at least one } i=1, 2, r$$

The *F* test statistic for the statistical significance of ρ_i^2 is:

$$F = \frac{1 - \lambda_i^{\frac{1}{r}} \frac{sd_2}{sd_1}}{\lambda_i^{\frac{1}{r}} \frac{sd_2}{sd_1}} \sim f_{sd_1, sd_2, \alpha}$$

Here:

$$\lambda_i = \prod_{s=1}^s (1 - r_i^2); s = \min(p, q); sd_1 = pq; sd_2 = wt - \frac{1}{2}pq + 1; w = n - \frac{1}{2}(p + q + 3); t = \sqrt{\frac{p^2q^2 - 4}{p^2q^2 - 5}}$$

Where:

N = The number of cases,

ρ = The number of variables in the X set,

q = The number of variables in the Y set,

r_i^2 : represents the eigenvalues of $\sum_1^{-1} \sum_2^{-1} \sum_2^2$ or the squared canonical correlations.

Statistical Analysis

All the computational work was accomplished using Statistical Package for Social Sciences statistical software version 26 IBM SPSS, 2019. Descriptive statistics such as means and standard deviation were analysed for a summary of the measured traits. Pearson's correlation was used for phenotypic correlation among measured traits. A Canonical correlation was used for all the canonical correlation analysis.

Results

The descriptive statistics for the examined traits at birth and weaning are shown in Table 1 while Table 2 shows a correlation among growth traits. Descriptive statistics indicated that WW mean value (15.2±22) was greater than the mean value of BW (3.77±1.0). Descriptive statistic for measured traits shows that all traits measured at weaning have greater mean value than all traits measured at birth. Correlation, highest correlations ($P < 0.01$) were estimated among HG and WH ($r = 0.82$) at birth; RH_3 and BL_3 ($r = 0.80$) at weaning; and WW and BW ($r = 0.80$) for interrelationships between birth and weaning. The lowest correlations ($P < 0.05$) were estimated among HW and BW ($r = 0.03$) at birth; WH_3 and RH_3 ($r = 0.22$) at weaning period; BL_3 and SH ($r = 0.04$) for interrelationships among birth and weaning, respectively.

Although studies show that birth time is essential indicators of weaning, it is challenging to explain the association among the traits simultaneously. For this cause, six canonical correlation coefficients were predicted to describe the interdependence among the studied variable set, since the number of canonical correlations that need to be explained is the lowest number of traits at birth or weaning period (Table 3). The likelihood ratio test shows a significant at first canonical correlation coefficient ($r = 0.881$) ($P < 0.01$). The results indicate the relation among the first canonical correlation variables (U_1 and V_1).

Canonical weights were specified for the first pair of canonical variables in Table 4.

The coefficients highlight the outcome that birth has on the weaning. On that account, the variables can be used to come up with defined optimal linear combinations with the use of standardized canonical coefficients as, $U_1 = (-0.97BW) - (0.05HG) + (0.04RH) - (0.15BL) + (0.07WH) - (0.06SH)$ $V_1 = (-1.00WW) - (0.22HG_3) - (0.30RH_3) + (0.46BL_3) - (0.14WH_3) + (0.09SH_3)$

Consistently, if the values of BW, HG, BL and SH reduce at birth, the readings of WW, HG_3, RH_3 and WH_3 at weaning will reduce, meanwhile the readings of the BL_3 and SH_3 at weaning will increase. Variables with higher canonical loadings given more to the multivariate

association among growth traits evaluated at birth and the weaning from non-descript goat kids (Table 5).

Withers height and sternum height were more influential informing V_1 compared to other characters and also in forming U_1 at birth according to loadings for the weaning time. Meanwhile, the cross-loadings suggest that WH_3 and WH gave the most to canonical variants V_1 and U_1 , respectively (Table 6).

First canonical variables U_1 explained 21.2% of the overall difference in the birth set, while 0.165 redundancy measures propose that about 16.5% of the ratio was explained by canonical variable V_1 . It was also found that the First canonical variables U_1 explained 29.5% of the overall difference in the weaning set, while 0.229 redundancy measures propose that about 16.5% of the ratio was described by canonical variable U_1 (Table7).

Table 1: Descriptive values for examined character (n = 50)

For Birth Period	X Variable Set Mean ± SD*	For Weaning Period	Y Variable Set Mean ± SD*
Birth Weight (BW)	3.7±1.0	Weaning Weight (WW)	15.2±2.2
Heart Girth (HG)	38.4±5.4	Heart Girth (HG_3)	56.5±4.4
Rump Height(RH)	36.0±4.1	Rump Height (RH_3)	46.5±4.6
Body Length (BL)	34.8±4.6	Body Length (BL_3)	52.6±4.6
Withers Height (WH)	35.8±4.0	Withers Height (WH_3)	47.2±2.8
Sternum Height(SH)	27.0±5.7	Sternum Height(SH_3)	33.3±5.5

*SD: Standard deviation

Table 2: Phenotypic correlation between traits weaning and birth

	BW	HG	RH	BL	WH	SH	WW	HG_3	RH_3	BL_3	WH_3
HG	0.21										
RH	0.24	0.48**									
BL	0.10	0.81**	0.32*								
WH	0.03	0.82**	0.47**	0.76**							
SH	0.04	0.76**	0.19	0.74**	0.81**						
WW	0.85**	0.28	0.21	0.18	0.09	0.64**					
HG_3	0.61**	0.18	0.10	0.08	0.04	0.26	0.54**				
RH_3	0.22	0.22	-0.02	0.28*	0.17	0.22	-0.02	0.34*			
BL_3	0.23	0.24	0.09	0.19	0.07	0.04	0.30*	0.54**	0.80**		
WH_3	0.06	0.18	0.14	-0.22	0.07	0.52**	0.54**	-0.02	0.22	0.28*	
SH_3	0.38**	-0.07	-0.07	0.13	-0.10	0.64**	-0.02	0.30*	-0.16	-0.03	0.32

BW = body weight, BL= body length, RH = rump height, WH = withers height, SH = sternum height, HG = heart girth.

** . Correlation is significant at the 0.01 level (1-tailed)

* . Correlation is significant at the 0.05 level (1-tailed)

Table 3: Summary results for the CCA

Pair of canonical	Canonical correlation	Square canonical	Eigenvalue	DF	Likelihood ratio	Probability Pr>F
U_1V_1	0.881	0.776	3.479	36	0.134	0.000
U_2V_2	0.448	0.200	0.251	25	0.602	0.663
U_3V_3	0.430	0.184	0.227	16	0.753	0.742
U_4V_4	0.212	0.044	0.047	9	0.925	0.950
U_5V_5	0.178	0.030	0.033	4	0.968	0.847
U_6V_6	0.021	0.000	0.000	1	1.000	0.893

Table 4: Standardized canonical coefficients for canonical variables

X – Variable set	Y – Variable set												
	BW	HG	RH	BL	WH	SH	WW	HG_3	RH_3	BL_3	WH_3	SH_3	
U_1	-0.97	-0.05	0.04	-0.15	0.07	-0.06	V_1	-1.00	-0.22	-0.30	0.46	-0.14	0.09

Table 5: Canonical loadings of the original variables with their canonical variables

X – Variable set	Y – Variable set												
	BW	HG	RH	BL	WH	SH	WW	HG_3	RH_3	BL_3	WH_3	SH_3	
U_1	-0.98	-0.34	-0.24	-0.27	-0.16	-0.18	V_{1s}	-0.97	-0.69	-0.31	-0.46	-0.08	-0.21

Table 6: Cross loading of the original variables with opposite canonical variables

X – Variable set	Y – Variable set					
	BW	HG	RH	BL	WH	SH
U ₁	-0.87	-0.30	-0.21	-0.24	-0.14	-0.16
Y – Variable set	X – Variable set					
	V ₁	WW	HG_3	RH_3	BL_3	WH_3
	-0.86	-0.61	-0.27	-0.40	-0.07	-0.19

Table 7: The explained total variation ratio by canonical variables for the variable sets

X-variable set	Y-variable set			
	Variance extracted	Redundancy	Variance extracted	Redundancy
U ₁	0.21	V ₁ 0.17	V ₁ 0.30	U ₁ 0.23

Discussion

The indigenous goats grow well under the harsh conditions and provide mainly milk, meat and cash income which is vital in the livelihood of rural people. The study firstly used Pearson’s correlation to examine the association among growth traits at birth and weaning. The findings suggest that HG had a correlation with WH at birth weight, while BL₃ and RH₃ had a correlation at weaning time. Secondly, a canonical correlation was used to examine the interdependence among birth time and weaning time. WW and BW have shown interrelations. The lowest correlations were predicted between HW and BW for birth time; RH₃ and WH₃ for weaning. SH and BL₃ for the interdependence between birth and weaning time. This study has revealed that WH and SH were factors that had more influence in this relation. This information on results will assist the breeders to select the best animals at birth to improve performance at weaning. These findings are in line with several studies (Çankaya and Kayaalp, 2007; Tahtali *et al.*, 2012; Ozen *et al.*, 2021). Çankaya and Kayaalp, (2007) estimated the association between live weight (y set) and some morphological traits in German sheep breeds using canonical correlation. The estimated canonical correlation coefficient was significant. Ozen *et al.* (2021) used canonical correlation analysis to study the association among seven biometric traits evaluated at weaning time (X-Set) and six months age (Y-Set) in a multidimensional form from an overall of seventy-one male lambs. The generated canonical dimensions had seven pairs and only the first pair was deemed statistically significant. The growth traits measures are important in the improvement of live weight.

Conclusion

The current paper has found the relationship between growth traits at birth and weaning time. Withers height, heart girth, body length and body weight were the most prominent factors in this relation. This information will help farmers in the selection of the best animal at birth and weaning to improve the bodyweight of the animal. More research needs to be performed on goats as the literature is minimal.

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Author’s Contributions

Mokoena Kwena: Fieldwork and draft of manuscript.

Thobela Louis Tyasi: Experimental design, data analysis, editing the manuscript and approved the final manuscript.

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