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Assessing the logistic growth function in connection to growth metrics

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Abstract

This study set out to establish which of the different non-linear growth curve model - Von-Bertalanffy, Gompertz, Richards, Weibull, exponential function, and logistic - was most suitable for characterising the growth curve. A total of 715 female and 340 male kids body weight records, taken every three months from birth to the twelfth month of life, were included. Males had a higher asymptotic live weight ("A") than females did. The maturity rate ("C") of males was higher than that of females. In conclusion, live weight as a function of age for male and female goats and sheep was best calculated using non-linear growth models. Calculations were made for residual SS (SSE), residual MS (MSE), AICC, and R². The various goods an animal produces, including milk, meat, and other foods, are impacted by its growth. This phase of the animal's existence is crucial. Since it is hard to interpret a series of weight-age data points analytically, it is preferable to analyse animal growth statistically. Understanding evolutionary change is essential to creating successful breeding strategies, and this is where growth curve inheritance comes in.

Keywords: Body weight, goat, sheep

Introduction

A variety of non-linear mathematical functions have been used to define the growth curve for evaluating response to a particular treatment at different times, interaction between and within populations for identifying heavier animals at an early age, and Gompertz (Laird, 1965) [8], Bertalanffy (Bertalanffy, 1957) [3], Logistic (Nelder, 1961) [11], and negative exponential (Bathaei and Leroy, 1998) [1] (Magotra *et al.*, 2021) [9].

Numerous studies have been undertaken in animal research to define the growth patterns of the animals. In addition to fitting standard models like simple linear or exponential growth to size-at-age data, these research also took into account models with sigmoidal (S-shaped) growth curves, the Brody model, and negative exponential growth models. The latter is modelled by Richards, von Bertalanffy, Gompertz, and Verhulst (logistic growth). A Google Scholar search yielded about 22,500 and 15,500 results, respectively, for articles on the application of the Brody model for sheep and goats. The Verhulst model gave approximately 5500 and 3500 hits, the Gompertz model produced roughly 4000 and 2000 hits, and the von Bertalanffy model produced roughly 2500 and 1500 hits (Brunner and Kühleitner, 2020) [5].

Materials and Methods

The sheep and goats were maintained on a vast field grazing method. The goats grazed on the pasture for six to eight hours every day. The trees, bushes, and grasses that are available for the goat are classified as follows: Monsoon (Kair, Dhaman, Dudh, Patharchatta, Motha, Akra, and

Thur), winter (Neem, Motha, Akra, Keekar, and Beri), and summer (post-harvest leftover residue of Gramme pea (Chickpea), Babul, Kair, and Khejri).

Statistical Analysis

Body weights were standardized for 30, 60, 120, 150, 210, 240, 300 and 330 days using the following methodology (Warwick and Legates, 1979) [21].

$$P_i = P_{near_i} + ADG (i - \text{age } P_{near_i})$$

where P_i is the standardized weight at standard age (i), P_{near_i} is the weight nearest to standard age (i), ADG is average daily gain considered among the weights after standard age (i) and before standard age (i), (i) is age to which weight is standardized, and age P_{near_i} , age to weight nearest to standard age (i) considering.

Average daily gain in the body weight of individual animal was calculated by using the following formula (Brody, 1964) [4].

$$\text{Average daily gain} = \frac{W_2 - W_1}{t_2 - t_1}$$

Where: W_2 = Final body weight (kg); W_1 = Initial body weight (kg); t_2 = Age of the animal at the end of the period (days); t_1 = Age of the animal at the beginning of period (days)

Different mathematical models were used to estimate growth curve parameters using Systatics 10 software.

Non-linear growth curve models	Equations
Logistic (Nelder, 1961) ^[11]	$W_t = A \times (1 + B \times e^{-kt})^{-1} + \varepsilon$

Where: W_t = the expected body weight (Kg) at 't' time; A= is the asymptotic weight; B= the folding point of growth; K = the rate of growth; m = Shape parameter; ε = random error; e = the base of natural logarithm; t = time (birth to 12th month of age).

Residuals were plotted graphically which gave an accuracy of the model to fit the growth curves.

$$e = y_i - \hat{y}_i$$

Where: y_i = Observed body weight at time "t"; \hat{y}_i = Predicted body weight by regression model at time "t".

The Mean Absolute Error (MAE), Mean Absolute Percentage Error (MAPE) was calculated as below (Topal and Balukbasi, 2008) ^[17].

$$MAE = \frac{\sum_i^n |y_i - \hat{y}_i|}{n}$$

$$MAPE = 1/n \sum_i^n \frac{|y_i - \hat{y}_i|}{y_i}$$

Where: y_i = Observed body weight at time "t"; \hat{y}_i = Predicted body weight by regression model at time "t"; n = Number of observations (data points).

In the case the sample size is smaller than the number of model parameter (NK<40), the AIC might not be accurate then after use of Akaike's information Criteria (AICC) was appropriate and calculated as using the equation Motulsky and Christopoulos (2004) ^[10].

$$AIC_c = AIC + \frac{2p(p+1)}{n-p-1}$$

Where: AIC = Akaike's Information Criteria.

Therefore, AICC is a good static for comparison of models of different complexity because it adjust the residual sum of squares (RSS) for number of parameters in the model. A smaller numerical value of AICC indicates a better fit when comparing models.

The Chi-square (χ^2) values is used to designate the relationship between actual and predicted body weights whether there is a significant difference between the predicted and the observed body weights.

$$\chi^2 = \frac{(y_i - \hat{y}_i)^2}{\hat{y}_i}$$

Where: Y_i = Observed body weight at time "t"; \hat{y}_i = Predicted body weight by regression model at time "t".

Results and Discussion

Male, female, and both sexes had estimated asymptotic mature live body weight parameters (A) of 22.49±0.88, 20.99±0.88, and 21.46±0.87 kg, respectively. The results were in close agreement with the reports of Yadav *et al.*, (2009) ^[22] as 20.60 Kg in both sexes of Marwari sheep whereas, In contrary the present result that lower parameter (A) was observed by Thirunavukkarasu *et al.*, (2017) ^[16] as 19.34, 16.76 and 17.50 Kg in male, female and both sexes of Mechari sheep, respectively.

The study by Ganesan *et al.* (2015) ^[7] reported higher estimates of parameter (A) as 36.44 and 25.99 kg in male and female Madras red sheep. On the other hand, Tsukahara *et al.* (2008) ^[18] reported higher parameter (A) for both sexes, with values of 27.00 kg in Kambing Katjang goat, Tatar *et al.* (2009) ^[15] reporting 32.03 kg in young hair goat, Paul *et al.* (2016) ^[23] reporting 32.53 kg in non-descript goat, and Waheed *et al.* (2016) ^[19] reporting 27.10 kg in Thalli sheep.

For the male, female, and both sexes, the Folding Point of Growth Curve (B) for the Logistic Growth Model was found to be 3.84±0.58, 3.87±0.63, and 3.89±0.62, respectively. The results of Paul *et al.* (2016) ^[23], which measured 3.75 in non-descript goats, and Thirunavukkarasu *et al.* (2017) ^[16], which measured 3.98 in Mechari sheep for both sexes, supported the current investigation.

The higher parameter (B) was found to be 4.38 in Kambing Katjang goat by Tsukahara *et al.* (2008) ^[18] and 4.04 in both sexes of Young hair goat by Tatar *et al.* (2009) ^[15]; on the other hand, Behzadi *et al.* (2014) ^[2] observed as 5.09 and 5.09 in Baluchi sheep and 4.05 and 4.06 in Mechari sheep for male and female by Thirunavukkarasu *et al.* (2017) ^[6].

For the male, female, and both sexes, the rates of maturing and growth rate (K) for the Logistic Growth Model were found to be 0.41±0.05, 0.41±0.05, and 0.42±0.05, respectively. The current result was consistent with results from Nimase *et al.* (2018) ^[13], showing 0.47 and 0.50 in Madgyal sheep, respectively, in the male and female.

Behzadi *et al.* (2014) ^[2] found lower estimates of parameter (K) for male and female Baluchi sheep, namely 0.02 and 0.02; Thirunavukkarasu *et al.* (2017) ^[16] found lower estimates, namely 0.01 and 0.01 for male, female, and both sexes of Mechari sheep, respectively.

Contrary to the current findings, Tatar *et al.* (2009) ^[15] reported a higher parameter (K) for both sexes in young hair goats, with a value of 0.66, while Waheed *et al.* (2016) ^[19] reported a value of 0.60 in Thalli sheep.

Conclusion

The study discovered that a variety of factors contribute to variation in growth curve model parameters, including but not limited to flock size, species, breed, selection techniques, environmental and managerial conditions, farmer socioeconomic position, and nutritional traits of breeds and species.

Declarations of Competing Interest

The authors declare that they have no conflicts of interest associated with this publication.

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