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Article

Variability in the Initial Growth Trajectory of Captive Bred Snakes of the Genus *Epicrates*

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Abstract: The goal of this study was to characterize the initial growth trajectory of *Epicrates assisi*, *Epicrates cenchria*, and *Epicrates crassus* snakes raised in captivity. An exponential model with parameters for birth weight (A) and growth rate (k) was fitted for each animal. There was a significant difference A and k among the three species, the means (standard error) for A and k parameters were 31 (0.3) g and 0.0027 (0.00006) for *E. assisi*, 27.6 (0.6) g and 0.0046 (0.00013) for *E. cenchria* and 25.9 (1.4) g and 0.0033 (0.00027) for *E. crassus*. We showed a significative variability between species for initial growth trajectory. In addition, we also demonstrated the existence of variability within species, which can be used to change the growth trajectory of snakes, being a tool for selection and genetic improvement.

Keywords: growth curve; performance; ex situ; rainbow boa; reproduction

1. Introduction

The *Epicrates* genus is composed of five species which are endemic at the neotropical region, occurring from Nicaragua to northwestern Argentina. In Brazil they are distributed across almost all biomes [15]. They are popularly known as “salamanta” or “rainbow boa” due to the iridescent effect of the dorsal scales under the incidence of light. All species of *Epicrates* are carnivorous constrictors with aglyphous dentition.

Snakes have increasingly been bred as pets around the world. Their diet in the wild consists of small mammals, amphibians, lizards, and birds. In captivity, their diet consists of mice for the smaller and younger snakes and rats for the larger ones [17]. Pet rainbow boas are usually sold once they are large enough to receive a transponder-type microchip with a universal identification number, which occurs between the third and fifth month of life.

The use of growth curves is important to understand the development of an animal both in captivity and in the wild. Fitting of nonlinear growth curves models allows to summarize a massive age-weight databases into a few parameters for studying and characterize the growth of the population; identify the superior animals at younger ages; evaluate the development of animals over time and study the interactions of responses of treatments with time [25]. Once genetic variability allows individuals to respond in different ways to changes that occur in the environment and are therefore, of great importance for the evolution of species.

Knowing the growth trajectory of these snakes, as well as the factors that influence their parameters, is important to predict the growth of animals in a population and explore the best management strategies. As sexual maturity is directly related to body size, low growth rate may result in reproductive delay of the animals. Therefore, the objective of this study was to analyze the variability and characterize the growth trajectory in the early phase of captive bred *Epicrates assisi*, *Epicrates cenchria*, and *Epicrates crassus*.

2. Materials and Methods

A database with repeated measurements of age-weight records of *Epicrates assisi*, *Epicrates cenchria* and *Epicrates crassus* snakes, born in captivity between 2017 and 2021, was provided by a commercial snake farm located in Betim, Minas Gerais (Brazil).

Animals were identified and weighed after birth and housed individually in clear plastic boxes measuring 22.2 x 15 x 6 cm with paper towel substrate. A bowl type water trough and an artificial heating source per box were used to provide a thermal gradient with a temperature range between 33°C to 35°C. The room had also had lighting (photoperiod 12:12 hours), temperature and humidity controlled. The animals were weighed every fifteen days on electronic scales accurate to one gram.

The first feeding of the offspring occurred after skin moulting, it was always used mouse (*Mus musculus*) neonates. Neonates of mice (*Rattus norvegicus*) could be offered at subsequent feedings. All animals were fed weekly at the rate of 5 to 10% of body weight and the water was provided ad libitum.

As this is an observational study with a database obtained under commercial conditions and uncontrolled variations, there was neither a planned experimental design nor the need for protocol to be approved by the Ethics Committee (CEUA).

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2.1. Data Editing

Only data from animals (females and males) *E. assisi*, *E. cenchria*, and *E. crassus* species born between the years 2017 and 2021, with at least five valid weightings, the first weighing being performed until the 30th day of life and the last weighing after 180 days of life were considered. After editing, the database was composed of 7210 weight data from six hundred and twenty animals, as presented in Table 1. For the individual trajectories of the animals see Figure 1.

Table 1. Descriptive statistics regarding the number of animals, weighings, and initial and final ages, according to species and sex.

Species	Sex	Animals	Starting age (days)			Final age (days)			Number of weighings			Total
			Med	Min	Max	Med	Min	Max	Med	Min	Max	
<i>E. assisi</i>	F	129	2.3	0	25	289.6	184	475	12	5	36	1555
<i>E. assisi</i>	M	138	1.9	0	18	281.1	184	419	12.5	5	36	1721
<i>E. cenchria</i>	F	141	1.9	0	21	255.9	182	425	11.72	5	30	1653
<i>E. cenchria</i>	M	131	1.5	0	15	255.7	190	436	10.9	6	44	1426
<i>E. crassus</i>	F	38	0.7	0	6	302.1	195	415	11.2	6	21	425
<i>E. crassus</i>	M	43	0.8	0	8	284.2	186	406	10	6	21	430

¹ Tables may have a footer.

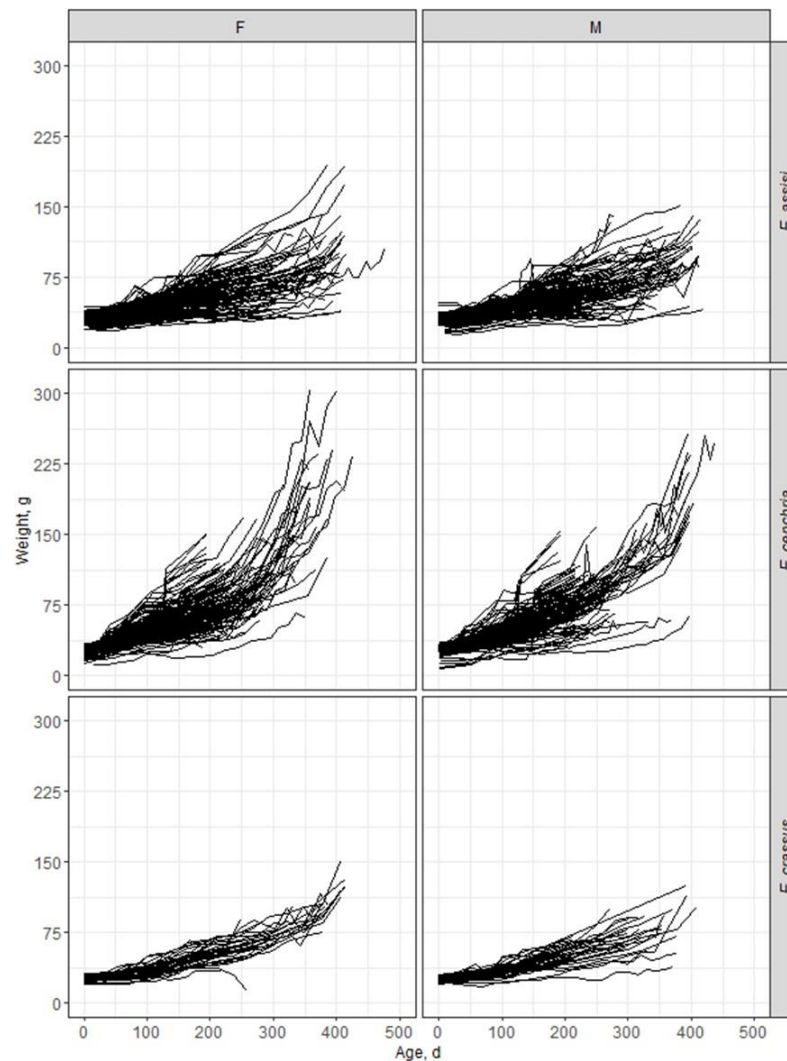


Figure 1. Figure 1. Individual growth trajectories observed for female (F) and male (M) snakes of the captive-bred species *Epicrates assisi*, *E. cenchria*, and *E. crassus*.

There were variations in relation to age at first weighing. From the years 2017 and 2019, the animals were weighed earlier while the animals born in 2020 and 2021 were weighed later, especially for *E. assisi* in 2021. Because the data was obtained from a commercial breeding farm, the variations were not controlled as well as management strategies varied over the seasons of birth which explains the differences in relation to age at first weighing of animals.

No specific pattern was observed for maximum age at weighing across years. However, the maximum age was higher for animals of the *E. crassus* species, intermediate for *E. assisi* and lower for *E. cenchria*, specifically in the years 2020 and 2021. As these animals are destined for sale, weight taking is closed on the eve of the animal leaving the farm. This sales order is also related to the popularization of each species. *E. cenchria* are more valuable and desired because they exhibit more intense colors and because of the number of rare color pattern variations obtained by the hatchery. Thus, animals of this species are sold faster than others. Estimates of the A_i and k_i parameters of an exponential model were obtained for each animal using the `nls_table` function of the `forestmangr` package [6] of the R software [23]. The exponential model used was as follows:

$$P_{ij} = A_i^{(k_i I_j)}$$

where:

P_{ij} represents the weight of animal i , at age j – expressed in grams;

A_i represents the weight of animal i , at age j – expressed in grams;

k_i represents the weight of animal i , at age j – expressed in grams;
 I_j represents the weight of animal i , at age j – expressed in grams;
 P_j represents the weight of animal i , at age j – expressed in grams;

The estimates of parameters $A(\hat{A}_i)$ and $k(\hat{k}_i)$ of each animal were used to predict their weights at the same ages they were measured $P(\hat{P}_{ij})$. Next, the following adjustment criteria were calculated for each animal:

$$\begin{aligned} \text{MAD}_i &= \frac{\sum_{j=1}^{J_i} |\hat{P}_{ij} - P_{ij}|}{J_i} \\ \text{MSE}_i &= \frac{\sum_{j=1}^{J_i} (\hat{P}_{ij} - P_{ij})^2}{J_i} \\ v_i &= 100 \times \frac{\sum_{j=1}^{J_i} \frac{\hat{P}_{ij} - P_{ij}}{P_{ij}}}{J_i} \\ r_i &= \frac{\text{Cov}(\hat{P}_{ij}, P_{ij})}{\sqrt{\text{Var}(\hat{P}_{ij}) \text{Var}(P_{ij})}} \end{aligned}$$

where:

MAD_i represents the mean of the absolute values of the deviations of animal i – expressed in grams;

J_i represents the total number of measurements of animal i ;

MSE_i represents the mean squared residues of animal i – expressed in grams²;

v_i represents the average of the adjustment bias for animal i – expressed in percentage;

r_i represents the correlation between predicted and observed values of animal i ;

$\text{Cov}(\hat{P}_{ij}, P_{ij})$ represents the covariance between predicted and observed values for the weight of animal i at age j ;

$\text{Var}(\hat{P}_{ij})$ represents the variance of the predicted weights of animal i at age j ;

$\text{Var}(P_{ij})$ represents the variance of the observed weights of animal i at age j ; and the other terms as defined above.

2.2. Analysis of Variance and Contrasts

The four fitting statistics, described above, and the estimates of the A and k parameters were considered as variables response and subjected to analyses of variance considering the following statistical model:

$$y_{ijkl} = \mu + E_i + S_j + A_k + ES_{ij} + EA_{ik} + SA_{jk} + e_{ijkl}$$

where:

y_{ijkl} represents the value of the response variable in species i , sex j , born in year k , and repetition l ;

μ represents a general constant present in all observations;

E_i represents the effect of species i ;

S_j represents the effect of gender j ;

A_k represents the effect of birth year k ;

ES_{ij} , EA_{ik} , SA_{jk} represent the effects of the double interactions of the factors described above; and e_{ijkl} represents the random error associated with each observation.

The null hypotheses for differences between means (for some factor of the statistical model, for example: species, sex, and year of birth) adjusted (for the other factors of the model) were tested by means of contrasts, using Tukey's method for p-value adjustment in multiple comparisons, using the options of the emmeans package [12] of the R software [23].

3. Results

Estimates of birth weight and relative growth rate (Table 2) along with analyzes of the adjusted individual growth trajectories for variations between and within species (Figure 2).

Table 2. Means (standard error in parentheses) of fit criteria and estimates of birth weight and relative growth rate of female (F) and male (M) snakes of the captive-bred species *Epicrates assisi*, *E. cenchria* and *E. crassus*.

Species	Sex	N	MAD (g)	MSE (g ²)	Bias (%)	Corr	A (g)	k (g)
<i>E. assisi</i>	F	129	3.175 (0.134)	18.940 (1.711)	0.857 (0.085)	0.918 (0.009)	30.974 (0.488)	0.00281 (0.00008)
<i>E. assisi</i>	M	138	3.069 (0.141)	19.083 (2.164)	0.879 (0.095)	0.926 (0.008)	30.697 (0.459)	0.00286 (0.00008)
<i>E. cenchria</i>	F	141	4.785 (0.203)	41.461 (3.290)	0.974 (0.208)	0.950 (0.006)	27.855 (0.504)	0.00486 (0.00012)
<i>E. cenchria</i>	M	131	4.178 (0.169)	33.327 (2.639)	1.229 (0.194)	0.956 (0.005)	27.029 (0.444)	0.00491 (0.00012)
<i>E. crassus</i>	F	38	3.251 (0.195)	18.511 (2.443)	0.628 (0.245)	0.929 (0.026)	25.072 (0.696)	0.00353 (0.00017)
<i>E. crassus</i>	M	43	2.635 (0.206)	13.216 (2.711)	0.421 (0.141)	0.959 (0.007)	24.855 (0.510)	0.00355 (0.00012)

¹MAD- mean of the absolute values of the deviations; MSE- mean of the squared residuals; Corr- correlation; A- birth weight; K- relative growth rate.

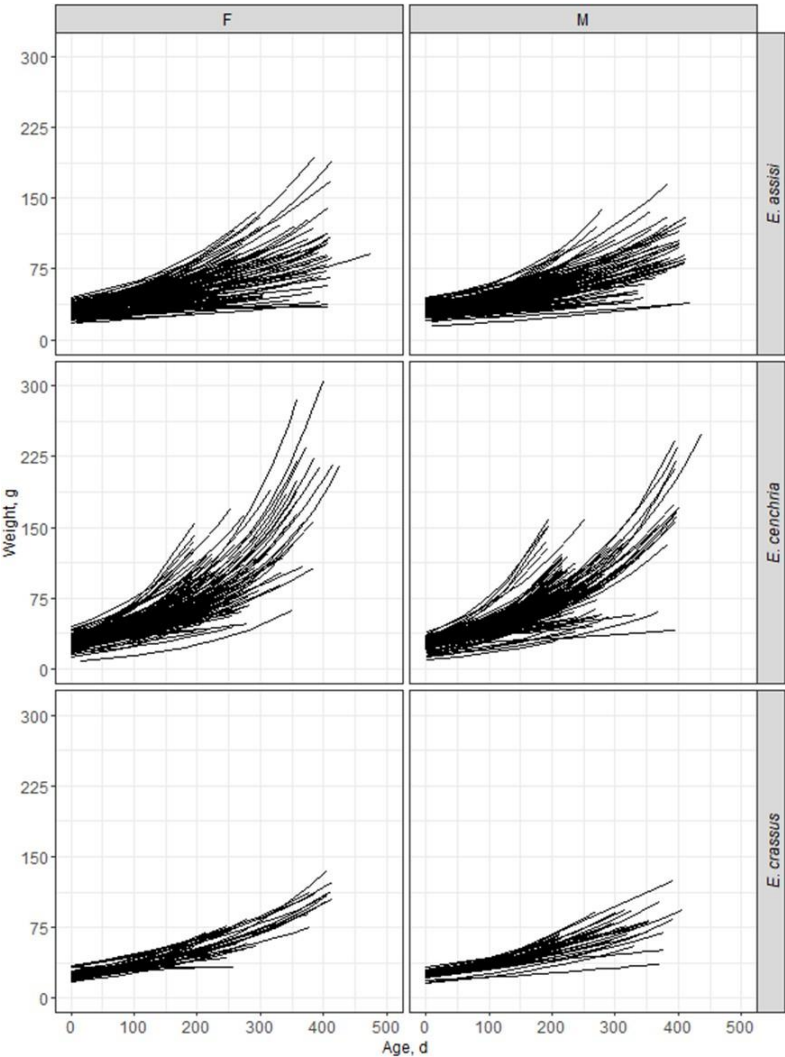


Figure 2. Individual growth trajectories fitted by an exponential model for female (F) and male (M) snakes of the captive-bred species *Epicrates assisi*, *E. cenchria*, and *E. crassus*.

The adjustment criteria indicated that the exponential model fit well with the snake growth trajectory. It was not possible to adjust other non-linear models, such as Gompertz, Richards, Von Bertalanffy among others, since the data analyzed contemplated only the exponential growth phase, being necessary to measure the weight for a longer period until a phase in which there was a decrease in growth rate and weight stabilization was identified.

The parameters indicating birth weight (A) and relative growth rate (k) were significantly influenced by species, year of birth, and the interaction of these factors (Table 3). The relative growth rate was also influenced by the interaction of sex and year of birth (Table 3).

Table 3. P-values associated with each source of variation of the statistical model used for the analysis of variance of the fit criteria¹ and the estimates of birth weight (A) and relative growth rate (k) of captive bred snakes of the species *Epicrates assisi*, *E. cenchria* and *E. crassus*.

Source of variation	MAD (g)	MSE (g ²)	Bias (%)	Correlation	A (g)	k (g)
Species (E)	0.0000	0.0000	0.0000	0.0006	0.0000	0.0000
Sex (S)	0.4487	0.5853	0.4321	0.4914	0.2816	0.4669
Year of birth (A)	0.0296	0.1281	0.7897	0.0000	0.0000	0.0000
E x S	0.3955	0.5999	0.2920	0.1940	0.7429	0.7746
E x A	0.0000	0.0000	0.0000	0.0145	0.0000	0.0000
S x A	0.5320	0.7264	0.2951	0.0253	0.2136	0.0283

¹MAD- mean of the absolute values of the deviations; MSE- mean of the squared residuals; Corr- correlation; A- birth weight; K- relative growth rate.

E. assisi was born heavier than *E. cenchria* and *E. crassus* (p<0.05). There was no difference (p>0.05) between birth weight of *E. cenchria* and *E. crassus*. The relative growth rate of *E. cenchria* was higher than the other two species (p<0.05) (Table 4).

Table 4. : Means (standard error in parentheses) of birth weight (A) and relative growth rate (k) for snakes of the species *Epicrates assisi*, *E. cenchria* and *E. crassus*, adjusted for the other effects included in the statistical model of the analysis of variance¹.

Species	A (g)	k (g)
<i>E. assisi</i>	31.0 (0.3) a	0.00277 (0.00006) a
<i>E. cenchria</i>	27.6 (0.6) b	0.00460 (0.00013) b
<i>E. crassus</i>	25.9 (1.4) b	0.00330 (0.00027) a

¹Means adjusted for the effects of sex, year of birth and by the double interactions involving the effects of species, sex and year of birth. Different letters in the same column indicate that the contrast between means is statistically significant (p<0.05), using Tukey’s method for p-value adjustment in multiple comparisons.

There was no significant difference for adjusted A in relation to the year of birth. In general, it is noticeable that the mean growth rates (k) decreased over time, however the only significant statistical differences (p<0.05) were in the years 2018 x 2019 and 2018 x 2021 (Table 5). The species x year of birth interaction was significant because there was a change in the magnitude of the year effect within species.

Table 5. Means (standard error in parentheses) of birth weight (A) and relative growth rate (k) for the year of birth for snakes of the species *Epicrates assisi*, *E. cenchria* and *E. crassus*, adjusted for the other effects included in the statistical model of the analysis variance¹.

Year of birth	A (g)	k (g)
2017	29.0 (1.3) a	0.00399 (0.00025) ab
2018	28.2 (0.4) a	0.00392 (0.00008) a

2019	28.2 (0.4) a	0.00356 (0.00008) b
2020	28.7 (1.8) a	0.00316 (0.00024) ab
2021	26.8 (1.2) a	0.00316 (0.00024) b

¹Means adjusted for the effects of sex, year of birth and by the double interactions involving the effects of species, sex, and year of birth. Different letters in the same column indicate that the contrast between means is statistically significant (p<0.05), using Tukey’s method for p-value adjustment in multiple comparisons.

4. Discussion

The parameters indicating birth weight (A) and relative growth rate (k) were significantly influenced by species, year of birth, and the interaction of these factors (Table 3). The relative growth rate was also influenced by the interaction of sex and year of birth (Table 3).

E. assisi was born heavier than *E. cenchria* and *E. crassus* (p<0.05). There was no difference (p>0.05) between birth weight of *E. cenchria* and *E. crassus*. The relative growth rate of *E. cenchria* was higher than the other two species (p<0.05) (Table 4).

There was no significant difference for adjusted A in relation to the year of birth. In general, it is noticeable that the mean growth rates (k) decreased over time, however the only significant statistical differences (p<0.05) were in the years 2018 x 2019 and 2018 x 2021 (Table 5). The species x year of birth interaction was significant because there was a change in the magnitude of the year effect within species.

The exponential model $y_i = A e^{ki}$ although not indicated for a long period of growth, is relatively common because of its simplicity. Growth can be easily described with only the initial and final weights [8]. Where A is the estimated initial weight; k is the relative growth rate and i is the age. Thus, following the development of *Epicrates* snakes after sexual maturity, after two to or three years, would be an excellent complement to this work.

Although *E. assisi* is endemic to the Caatinga biome (from the states of Piauí to southern Bahia and northern Minas Gerais in Brazil) it may occur in syntopy with *E. crassus* in southwestern Bahia state to northern Minas Gerais in transition areas between Caatinga and Cerrado [15]. Such characteristics may explain the similarity in growth profile between the species.

E. assisi had a higher birth weight in relation to the other species. This fact may be an adaptation to the conditions of their natural habitat, which is characterized by extreme droughts and dry periods typical of the semi-arid climate. Heavier offspring may have more chances of survival even with the lower availability of food.

E. cenchria inhabits the Amazon rainforest, considered to have the greatest biodiversity on the planet. The abundance of food and the favorable climate optimizes the development of this species, which making it one of the largest and heaviest snakes of the *Epicrates* family, possibly exceeding 2.20 meters.

The Cerrado biome presents an abundance of prey intermediate between Caatinga and Amazon rainforest. This fact may bring the ‘Cerrado salamanta’ closer to both the Caatinga and Amazon ‘salamanta’ being found similarities in birth weight between *E. crassus* and *E. cenchria*. Variations were detected regarding the individual growth trajectory and relative growth rate between and within species, these differences may have occurred because growth patterns vary, and the model used was not robust enough to fit all growth patterns in the same way. However, variability can be caused by some not identified difference in management or due to genetic differences between animals of the same species, which can be a meaningful material for the evolution of species. To better understand the variability, it would be necessary to have proper design in which the family of the parents of these animals or the molecular markers are known.

In commercial breeding, knowing the growth behavior of animals from birth is fundamental to the success of the activity. Animals that grow faster may be sold sooner, contributing to the reduction of operational and production costs. In the reproductive bias, fertility tends to increase as alongside the growth of the body of female snakes [?]. Size at birth can be influenced by female size and litter size [11]. Sexual maturity in these species is more related to body weight and size than age.

Thus, selecting snakes with higher growth rates until puberty becomes an important tool for genetic selection and herd improvement.

Growth rates among species can be totally different even among species whose adults are similar in size. This difference occurs due to physiological reasons that allow a higher rate of food conversion into mass [24].

Snakes continues to growth even after reaching reproductive status, females accumulate body reserves for long periods before reproduction to effect vitellogenesis. Reproductive performance and body weight may interact and vary according to environmental conditions and food availability [1]. In addition, larger females tend to give birth to larger offspring, which is desirable for the production system.

Some snake taxa exhibit sexual dimorphism by size. Snakes that exhibit male-male combat behavior have larger male individuals than females, while females of highly fertile species are larger than males [23]; [20]; [9]).[21] maintained a litter of *Bothrops fonsecai* under a controlled feeding regime and showed that females, even though they received the same amount of food as males, showed higher growth rates in all measured parameters.

The authors discuss that sexual dimorphism is independent of the amount of food ingested, but rather connected to the different food conversion between the sexes and suggest that this difference may be linked to the efficiency of energy assimilation by the female. A population of *Agkistrodon piscivorus* (Lacépède, 1789) was studied for 14 years by Ford (2002) and similar growth rates were found between the sexes at the juvenile stage. However, males grew more after reaching sexual maturity.

5. Conclusions

The present study provided evidence of the existence of significant variability for the initial growth trajectory. *E. assisi* is heavier than *E. cenchria* and *E. crassus* at birth, but the growth rate is higher in *E. cenchria* than in the other two species. The variability within species can be used to alter the growth trajectory of snakes, being a tool for selection and genetic improvement, since it is to elucidate growth patterns, the relationship between size and age, or even possible to identify the existence of differences between individuals of the same species.

6. Patents

This section is not mandatory, but may be added if there are patents resulting from the work reported in this manuscript.

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Data Availability Statement: We encourage all authors of articles published in MDPI journals to share their research data. In this section, please provide details regarding where data supporting reported results can be found, including links to publicly archived datasets analyzed or generated during the study. Where no new data were created, or where data is unavailable due to privacy or ethical restrictions, a statement is still required. Suggested Data Availability Statements are available in section “MDPI Research Data Policies” at <https://www.mdpi.com/ethics>.

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Conflicts of Interest: The authors declare no conflict of interest.

Sample Availability: Samples of the compounds ... are available from the authors.

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