

Non-linear curve adjustments widen biological interpretation of relative growth analyses of the clam *Tivela mactroides* (Bivalvia, Veneridae)

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Evaluation of relative (allometric) growth provides useful information to understand the development of organisms, as well as to aid in the management of fishery-exploited species. Usually, relative growth analyses use classical models such as the linear equation or the power function (allometric equation). However, these methods do not consider discontinuities in growth and may mask important biological information. As an alternative to overcome poor results and misleading interpretations, recent studies have suggested the use of more complex models, such as non-linear regressions, in conjunction with a model selection approach. Here, we tested differences in the performance of diverse models (simple linear regression, power function, and polynomial models) to assess the relative growth of the trigonal clam *Tivela mactroides*, an important fishing resource along the South American coast. Regressions were employed to relate parameters of the shell (length (L), width (W), height (H) and weight (SW)) among each other and with soft parts of the organism (dry weight (DW) and ash-free dry weight ($ASDW$)). Then, model selection was performed using the information theory and multi-model inference approach. The power function was more suitable to describe the relationships involving shell parameters and soft parts weight parameters (i.e. L vs. SW , DW , and $AFDW$, and SW vs. DW). However, it failed in unveiling changes in the morphometric relationships between shell parameters (i.e. L vs. W and H ; W vs. H) over time, which were better described by polynomial functions. Linear models, in turn, were not selected for any relationship. Overall, our results show that more complex models (in this study polynomial functions) can unveil changes in growth related to modifications in environmental features or physiology. Therefore, we suggest that classical and more complex models should be combined in future studies of allometric growth of molluscs.

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ABSTRACT

Evaluation of relative (allometric) growth provides useful information to understand the development of organisms, as well as to aid in the management of fishery-exploited species. Usually, relative growth analyses use classical models such as the linear equation or the power function (allometric equation). However, these methods do not consider discontinuities in growth and may mask important biological information. As an alternative to overcome poor results and misleading interpretations, recent studies have suggested the use of more complex models, such as non-linear regressions, in conjunction with a model selection approach. Here, we tested differences in the performance of diverse models (simple linear regression, power function, and polynomial models) to assess the relative growth of the trigonal clam *Tivela mactroides*, an important fishing resource along the South American coast. Regressions were employed to relate parameters of the shell (length (L), width (W), height (H) and weight (SW)) among each other and with soft parts of the organism (dry weight (DW) and ash-free dry weight ($AFDW$)). Then, model selection was performed using the information theory and multi-model inference approach. The power function was more suitable to describe the relationships involving shell parameters and soft parts weight parameters (i.e. L vs. SW , DW , and $AFDW$, and SW vs. DW). However, it failed in unveiling changes in the morphometric relationships between shell parameters (i.e. L vs. W and H ; W vs. H) over time, which were better described by polynomial functions. Linear models, in turn, were not selected for any relationship. Overall, our results show that more complex models (in this study polynomial functions) can unveil changes in growth related to modifications in environmental features or physiology. Therefore, we suggest that classical and more complex models should be combined in future studies of allometric growth of molluscs.

1. INTRODUCTION

Relative or allometric growth analysis allows a detailed evaluation of the proportionality among the body parameters of individuals, and is useful to estimate the production and biomass of a population from a single parameter such as length or width (Rainer 1985; Urban & Campos 1994; Vasconcelos et al. 2018). It also provides a better comprehension of modifications in life strategies of species unraveling important steps in their development (Katsanevakis et al. 2006; Rabaoui et al. 2007). Changes in growth rates are usually linked to changes in environmental features (e.g. food availability), physiology (e.g. gonad maturation and spawning periods), or biotic interactions (e.g. increase of competition or predation), and must be considered when evaluating morphological modifications during the ontogeny of the individuals (Rabaoui et al. 2007; Caill-Milly et al. 2014). The analysis of allometric growth is therefore an important tool to support exploitation and management of harvested species (e.g. improving the size-selectivity of fishing gears) (Vasconcelos et al. 2018), and essential for the proposal of effective measures to protect living resources (e.g., monitoring of stocks and limiting harvesting practices according to size-based analyses of communities) (Robinson et al. 2010).

Traditionally, allometric growth analyses are performed using the linear equation and power function (allometric equation) (e.g., Gaspar et al. 2001; 2002; MacCord & Amaral 2005; Vasconcelos et al. 2018). While the former ($y = a + bx$) corresponds to an equivalent increase in size of a body part y (dependent variable) and another body part x (independent variable; the reference dimension), the power function (allometric equation) determines the relationship between two parts of the body through an exponential equation $y = ax^b$, where b is a measure of the constant difference in the growth rates of the body parts x and y (Katsanevakis et al. 2006). However, the allometric exponent b is not necessarily constant and may exhibit breakpoints (i.e.

points of discontinuity in slope) for instance resulting from marked changes in the environment or physiology (Katsanevakis et al. 2006). As an alternative to overcome poor and misleading interpretation of results, more complex models that consider that nonlinearity and breakpoints may exist in the relationship of body parts (e.g., polynomial functions such as quadratic or cubic models) have been recommended (Hendriks et al. 2012; Katsanevakis et al. 2006). Preferably, researchers should consider a set of pre-established models and use model selection methods, such as the information theory and multi-model inference approach (Burnham & Anderson 2002b), to ascertain which better fits the data (Katsanevakis et al. 2006; Rabaoui et al. 2007).

Regarding economic important marine species, allometric growth analyses are relatively common for fishes, but much scarcer for benthic invertebrates (Vasconcelos et al. 2018). For bivalves, the growth and shape of the shell are normally influenced by both environmental (e.g., temperature, depth, currents, wave exposure, and sediment) and biological factors (predation, growth, and burrowing abilities) (see revision by Gaspar et al. 2002). Therefore, variation in these conditions may produce varying growth patterns of the shell and soft parts of the animals. An evaluation of the allometric growth of the bivalve *Pinna nobilis*, for example, registered significant variation in growth among five populations related to various environmental factors (Rabaoui et al. 2007). Marked changes in bivalve allometric growth are also expected to be associated with the maturity of individuals (i.e., the onset of reproduction), given that more energy is allocated to soft parts (mainly to the development and maturation of gonads) than to shell deposition (Bayne & Worrall 1980; Rabaoui et al. 2007). To better understand the development of key species and provide critical information for fisheries stock assessment and management, it is therefore essential that allometric growth studies are performed on a wider diversity of organisms and use a more comprehensive set of statistical models. Only through this

approach, growth variation among species, and between and within populations (i.e., spatial and temporal comparisons) can be unveiled.

In this study, we tested whether different models are equally adequate to estimate the allometric growth of the trigonal clam *Tivela mactroides* (Born, 1778), an important fishing resource in the Southeastern American coast (Denadai et al. 2005; Turra et al. 2016). Specifically, we compared the suitability of models traditionally used in allometric growth analyses (i.e., linear and power function models) with more complex models (i.e., polynomial functions), and investigated whether alternative models may lead to a better interpretation of the data and provide additional biological information about this species.

2. MATERIALS AND METHODS

2.1. *Model species*

The trigonal clam *T. mactroides* is a widespread species found along the South America coast, from Colombia to southeastern Brazil (Turra et al. 2014). This bivalve occurs from the shallow subtidal to the upper intertidal zone (Denadai et al. 2005), and is a main feeding resource for several species of fishes, sea-stars, and crabs (Turra et al. 2015a). Moreover, *T. mactroides* is economically important and intensively exploited by fishermen and recreational harvesters in many parts of South America (Turra et al. 2016).

2.2. *Study area*

This work was performed at Caraguatatuba Bay, which is located in southeastern Brazil (Fig. 1) and comprises several sandy beaches along a 16 km beach arch (Denadai et al. 2013). Wave energy is moderate at Caraguatatuba Bay because of the shadowing effect of São

Sebastião Island (Denadai et al 2005), but beach characteristics vary in a north–south orientation. The southernmost part has a wide intertidal ultradissipative terrace (800 m) with well-sorted fine sand. The northern part has a more heterogeneous slope with low tide dissipative terrace and fine poorly sorted sand (Denadai et al. 2005, Turra et al. 2014). The mean seawater temperature in Caraguatatuba Bay ranges from 19 °C in winter to 26° C in summer (Corte et al. 2015). Salinity is usually above 30 and significant variations only occur near the rivers that flow into the bay (Amaral and Nallin 2011).

2.3. *Sampling and laboratory procedures*

Field work was done following procedures described in Turra et al. (2015). Briefly, clams were sampled monthly from January 2003 until October 2004 in two areas with high abundance of *T. mactroides* (Denadai et al. 2005) (Fig. 1). The first southern area extended from Porto Novo to Palmeiras; and the northern area from Indaiá to Centro beach (Fig. 1). Both areas had 2000 m length and sampling was performed in the intertidal and subtidal zone of each area following different procedures.

In the intertidal zone of each area, sampling was performed at eight randomly sorted transects from 200 possibilities (i.e. the linear 2000 m length divided into 10-m intervals). Six or seven samples (squares of 0.5 x 0.5 m) were excavated to a depth of 10 cm at regular intervals in each transect. The number of samples collected at each transect depended on their length. A total of 120 samples (30 m²) were collected per month in the intertidal area.

In the subtidal zone, sampling was performed using a fishing boat at five different depths, i.e., 400, 800, 1200, 1600 and 2000 m, from mean low water (MLW, 0.0 m). At each depth, one 50-m dredging was performed at three randomly sorted transects perpendicular to the coast.

Thus, 30 samples were dredged per month (2 areas x 5 depths x 3 transects). Samples were taken using a rectangular dredge (70 x 25 cm) with 3.0 mm internal mesh size.

The sediment sampled at both intertidal and subtidal zones was washed with seawater over a 3.0-mm-mesh sieve, using buckets. Clams from both zones were collected, counted and measured for shell length to the nearest 0.01 mm with a digital caliper to examine the population dynamics of *T. mactroides* at the study area as reported in Turra et al. (2014). Almost all the individuals were returned alive to the sea after the measurements; however, approximately five individuals from each of the 37 size classes identified in this population (from 3.0 to 39.0 mm; 1 mm intervals) were taken for morphometric analyses, totaling 187 individuals. These specimens were maintained in clean, aerated seawater for 24 h to eliminate the feces. After this period, the soft parts were separated from the shells and dried at 60°C to achieve a constant weight and afterwards record the shell weight and the soft parts dry weight. Dry weight was used to avoid discrepancy in the amount of water retained or fluctuation in soft-parts weight due to changes in the physiological processes (Vasconcelos et al. 2008; 2018). Ash weight was obtained after incinerating the dried soft parts in a muffle furnace at 550°C for 5 h. The ash-free dry weight (organic content) was calculated by subtracting the ash weight from the dry weight. Thus, all individuals had the shell length (*L*), height (*H*), and width (*W*) measured (Fig. 2), and the shell weight (*SW*), soft parts dry weight (*DW*), and ash-free dry weight (*AFDW*) recorded.

2.4. Data analysis

The allometric growth of *T. mactroides* was analyzed using models that relate the shell length (independent variable) to the shell height, shell width, shell weight, soft parts dry weight, and ash-free dry weight (*L* vs *H*, *W*, *SW*, *DW* and *AFDW*). The relationships between shell width

(independent variable) and shell height (W vs H), and between the soft parts dry weight (independent variable) and the shell weight and ash-free dry weight were also evaluated (DW vs SW and $AFDW$).

We used five models for the comparisons (Table 1): (1) simple linear regression ($y = bx$), (2) second-order polynomial function ($y = bx + cx^2$), (3) third-order polynomial function ($y = bx + cx^2 + dx^3$), (4) fourth-order polynomial function ($y = bx + cx^2 + dx^3 + ex^4$), and (5) power function (allometric equation, $y = ax^b$). The candidate models for allometric growth were fitted to raw data and the fitted equations were compared based on the Kullback–Leibler (K-L) information theory and multi-model inference (MMI) (Burnham & Anderson 2002). The Kullback–Leibler (K-L) information theory can be interpreted as the distance from the approximating model to full reality, and minimization of K-L distance is essential for model selection (Katsanevakis et al. 2006). Multi-model inference, in turn, is a procedure where parameters are estimated from several different models rather than from one *a priori* selected model (Burnham & Anderson 2002).

We used the small-sample bias-corrected form of the Akaike Information Criteria, AIC_c (Hurvich & Tsai 1989) of the AIC (Burnham & Anderson 2002a) for the model selection, according to the equation:

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

where AIC is given by:

$$AIC = n \left(\log \left(2\pi \frac{RSS}{n} \right) + 1 \right) + 2k$$

where RSS is the residual sum of squares of the regression, n is the number of observations, and k is the number of estimated parameters of the regression plus 1. The model with the smallest AIC_c

value ($AIC_{c,min}$) was selected as the ‘best’ among the tested models. The AIC_c differences between adjusted functions, $\Delta_i = AIC_{c,i} - AIC_{c,min}$, were computed over all pairs of candidate models. According to Burnham & Anderson (2002), models with $\Delta_i < 2$ have substantial support, models with $4 < \Delta_i < 7$ have considerably less support, and models with $\Delta_i > 10$ essentially have no support and can be ignored. In the present work, we considered that all values of $\Delta_i < 2$ have substantial support for the model, and in the case of multiple models having $\Delta_i < 2$ we opted for the simplest model (principle of parsimony), i.e. the model with fewer estimated parameters.

2.5. Field Study Permissions

All activities complied with the license issued by the appropriate federal environmental agency (Ministério do Meio Ambiente (MMA)—Instituto Chico Mendes de Conservação da Biodiversidade (ICMBio) No. 19887-1; acronyms for, in English: Ministry of the Environment—Chico Mendes Biodiversity Conservation Institute)

3. RESULTS

High determination coefficients ($r^2 > 0.842$) were obtained in all models fitted to the relationships between the morphometric parameters of *T. mactroides* (Table 2). For most relationships, more than one model was suitable since they had values of $\Delta_i \leq 2$ (Table 2). The third-order polynomial suitably described all morphometric relationships excluding the L/W relationship. Nevertheless, considering the parsimony principle, the traditional power function was overall the most adequate model and better described five morphometric relationships (L/SW , L/DW , $L/AFDW$, SW/DW , and $DW/AFDW$). The third-order polynomial model was the

best descriptor for two relationships (L/H and W/H), whereas the fourth-order polynomial was selected as the best descriptor of the relationships L/W .

The relative growth between shell length and height was approximately isometric up to 20 mm L; above this size, shell length increased faster compared to shell height (Fig. 3). The relative growth between shell length and width was always positive allometric (i.e., length increased faster), but this relationship was stronger after reaching maturity (approximately 20 mm in L, Pietro 1980). The relationships between shell length and soft parts showed that the increase in soft parts (DW and $AFDW$) was slow in young individuals (below 20 mm L); followed by a rapid increase in soft parts weight was observed after reaching maturity. A similar relationship was observed between shell length and shell weight. The allometric growth between width and height oscillated during growth, mainly between 15 and 25 mm in shell width. The shell weight increased continuously (practically linearly) with soft parts dry weight. Similar relationship was established between ash-free dry weight and soft parts dry weight.

4. DISCUSSION

The usual approach when studying allometric growth in marine species is to *a priori* adopt the classical linear or allometric model (power function), which may have implications in the accuracy and precision of the estimated parameters (Katsanevakis et al. 2006; Rabaoui et al. 2007). When only these classical models are used, less informative conclusions may be reached by smoothing the real growth pattern (Protopapas 2007; Rabaoui et al. 2007). In this study, we used the information theory and multi-model inference approach (Burnham & Anderson 2002a) to compare the suitability of classical (i.e., linear and power function models) and more complex models (i.e., polynomial functions) to describe the allometric growth of *T. mactroides*. Our

results showed that classical models may adequately describe most morphometric relationships of this species. However, their use alone may hide subtle changes in the allometric growth and prevent a full understanding of growth patterns.

The power function (allometric equation) was the best model describing five of the eight morphometric relationships analyzed (mainly relationships involving the weight parameters such as shell weight, dry weight and ash-free dry weight). This function is commonly used in morphometric analyses of bivalves (e.g., Clasing et al. 1994; Urban & Campos 1994; Gaspar et al. 2001; Lomovasky et al. 2005), but is normally considered the weakest adjustment for many morphometric relationships in some marine invertebrates (Katsanevakis 2007; Rabaoui et al. 2007; Garraffoni et al. 2010). A main drawback in the use of the power function is that it does not consider breakpoints (i.e., marked changes in the allometric growth) and may hide major indications of morphological variability among individuals of different sizes (Rabaoui et al. 2007). Our results show that the power function best characterized the allometric growth between shell parameters and soft tissues, indicating that shell and somatic parts of *T. mactroides* grow with a constant allometric exponent, whereas polynomial models better described relationships between shell parameters. Likewise, Trussel (2000) found that changes in shell morphology in *Littorina* snails were not related to soft tissue variations, and McKinney et al. (2004) reported that allometric regression models best described changes in soft tissue content with shell length for 10 different species, including the ribbed mussel *Geukensia demisa*, the blue mussel *Mytilus edulis*, and the quahog *Mercenaria mercenaria*.

Several factors may influence the growth of soft tissue relative to shell parameters. Food availability, for instance, can strongly affect growth, storage and utilization of body reserves; thereby altering allometric relationships (McKinney et al. 2004). Similarly, temperature and

salinity may alter metabolic rates and determine net growth efficiency in bivalve species (Resgalla Jr. et al. 2007, Xiao et al. 2014). The constant allometric growth of shell measurements and weighings of *T. mactroides* at Caraguatatuba Bay is likely related to constant food supply to the clams due to the presence of three rivers (i.e., Santo Antônio, Lagoa and Juqueriquerê – Fig. 1) that flow directly into the bay and provide a large amount of suspended organic material (Corte 2015; Turra et al. 2015b). The absence of marked fluctuation in seawater temperature and salinity at the study area (Amaral and Nallin 2011, Corte et al. 2015) probably also contributes to the constant growth of soft tissues.

Although the power function was more precise to describe the morphometric relationship between shell parameters and somatic parts, it was not enough to unveil changes in the morphometric relationships between shell parameters. By adjusting polynomial functions, we were able to detect differences in morphometric relationships of shell parameters over the lifespan of *T. mactroides*. Changes in bivalves morphometric relationships are usually related to the maintenance of an area/volume ratio that is physiologically suitable for the prevailing environmental conditions (Gaspar et al. 2002; Rhoads & Pannella 1970). For instance, the shell of the pill clam *Pisidium subtruncatum* (Malm 1855) change from a rounded and thin-shelled form to an angular and thick-shelled form depending on the proportion of sand in the sediment (Funk & Reckendorfer 2008), which enables the individuals to burrow easier in coarse sediments. Similar modifications have also been observed in species that change their habits from active burrowing juveniles to sedentary and deeper burrowing adults (Statzner & Holm 1982), such as the razor clam *Ensis macha* that shows an abrupt change in shell shape during growth, from squarer to more elongated (Barón et al. 2004). This elongated form enables razor

clams to avoid predators by burrowing deeper with low energy requirements (Trueman 1967; Urban 1994).

Modifications in the allometric relationships between shell parameters of *T. mactroides* seem related to habitat changes and maturity of individuals. We observed that shell width and height had a reduction in their proportional growth in relation to shell length when individuals reached a length between 15 and 25 mm. This change is appropriate for the subtidal-intertidal migration undertaken by this species (Denadai et al. 2005). The smallest individuals are usually found in mud sediments in the subtidal level, where burrowing with a more-inflated shell is easier. As individuals reach maturity, they passively migrate to firm intertidal sandy substrata and their shell becomes longer and more compressed, potentially facilitating burrowing in this type of sediment. Maturity is also a frequent cause of a distinct change in morphology, given that more energy is allocated to the development of gonads than to shell growth (Bayne & Worrall 1980; Katsanevakis et al. 2006).

Besides providing further information for understanding the life history of species, the knowledge of allometric growth has implications for economic exploitation of fishing resources. In bivalves, the morphometric relationships between shell size and soft parts can help defining an optimum size for exploitation. For *T. mactroides*, the analyses of allometric growth indicate that more energy is invested in shell growth during early phases of the life cycle, thereby increasing the protection and survival of juveniles. Only after reaching maturity, a faster increase in somatic tissues was recorded. Thus, harvesting of *T. mactroides* should preferably target individuals with shell length between 25-30 mm. This would ensure that individuals are already able to reproduce, thus providing a better income to local harvesters since a higher meat yield (ratio meat weight/shell weight) would be achieved.

5. CONCLUSIONS

Our results show that the classical power function is useful to evaluate the relative growth of *Tivela mactroides*. Nevertheless, the use of this model alone may hide subtle changes in morphology related to environmental or physiological processes such as habitat changes and maturity of individuals. Only by applying more complex models, in the present case polynomial models, we perceived subtle modifications in the allometric growth of *T. mactroides*. In this regard, we reinforce the recommendation that allometric growth analyses should consider a set of pre-established models including traditional and more complex models. Certainly, this would improve allometric analyses and provide stronger and more informative conclusions.

6. ACKNOWLEDGEMENTS

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Table 1(on next page)

Models and their respective equations and number of parameters estimated plus one (k). All models consider origin = 0.

- 1 Table 1. Models and their respective equations and number of parameters estimated plus one (k).
- 2 All models consider origin = 0.

Model	Equation	k
Simple regression (Ln)	$y = bx$	2
Second-order polynomial (Qd)	$y = bx+cx^2$	3
Third-order polynomial (Cb)	$y = bx+cx^2+dx^3$	4
Fourth-order polynomial (Qt)	$y = bx+cx^2+dx^3+ex^4$	5
Power-function (Pf)	$y = ax^b$	3

3

Table 2 (on next page)

Tivela mactroides.

Summary and comparison among the determination coefficient (r^2), standard error of the estimate (SE), residual sum of squares (RSS), Akaike information criteria corrected for small samples (AIC_c), and difference between AIC_c (Δ_i) of all models.

L, shell length; W, shell width; H, shell height; SW, shell weight; DW, soft parts dry weight; AFDW, ash-free dry weight. Ln, linear model; Qd, second-order polynomial model; Cub, third-order polynomial model; Qt, fourth-order polynomial model; Pf, power function. Most suitable model according to AIC_c and parsimony principle are highlighted in bold. When $\Delta_i < 2$, the model with the smallest number of parameters was selected.

Table 2. *Tivela mactroides*. Summary and comparison among the determination coefficient (r^2), standard error of the estimate (SE), residual sum of squares (RSS), Akaike information criteria corrected for small samples (AIC_c), and difference between AIC_c (Δi) of all models. L, shell length; W, shell width; H, shell height; SW, shell weight; DW, soft parts dry weight; AFDW, ash-free dry weight. Ln, linear model; Qd, second-order polynomial model; Cub, third-order polynomial model; Qt, fourth-order polynomial model; Pf, power function. Most suitable models according to AIC_c and parsimony principle are highlighted in bold. When $\Delta i < 2$, the model with the smallest number of parameters was selected.

Relation	Model	r^2	SE	RSS	AIC_c	Δi
L/W	Ln	0.997	0.857	136.871	476.392	94.070
	Qd	0.998	0.753	104.985	428.861	46.539
	Cub	0.998	0.689	87.519	396.923	14.602
	Qt	0.998	0.661	80.037	382.322	0.000
	Pf	0.985	0.799	118.223	451.068	68.747
L/H	Ln	0.999	0.730	99.106	416.019	107.264
	Qd	0.999	0.574	61.111	327.669	18.914
	Cub	0.999	0.546	55.005	310.073	1.318
	Qt	0.999	0.543	54.005	308.755	0.000
	Pf	0.996	0.617	70.424	354.194	45.439
L/SW	Ln	0.861	1.730	556.922	738.824	464.003
	Qd	0.987	0.530	52.115	297.891	23.070
	Cub	0.989	0.499	45.964	276.496	1.675
	Qt	0.989	0.501	45.942	278.516	3.695
	Pf	0.976	0.499	46.066	274.821	0.000
L/DW	Ln	0.844	0.094	1.637	-342.574	250.325
	Qd	0.959	0.048	0.429	-586.848	6.052
	Cub	0.960	0.047	0.415	-591.088	1.811
	Qt	0.961	0.047	0.411	-590.689	2.210
	Pf	0.918	0.047	0.415	-592.899	0.000
L/AFDW	Ln	0.842	0.086	1.351	-374.856	251.689
	Qd	0.959	0.044	0.351	-619.540	7.004
	Cub	0.961	0.043	0.337	-624.709	1.836
	Qt	0.961	0.043	0.334	-624.112	2.433
	Pf	0.918	0.043	0.338	-626.544	0.000
W/H	Ln	0.999	0.835	62.982	331.243	20.198
	Qd	0.999	0.837	61.612	329.198	18.153
	Cub	0.999	0.830	55.291	311.045	0.000
	Qt	0.999	0.7651	107.127	436.837	125.792
	Pf	0.992	0.8365	129.446	468.027	156.982
DW/SW	Ln	0.962	0.047	0.399	-597.694	0.266
	Qd	0.962	0.047	0.395	-597.676	0.285
	Cub	0.963	0.046	0.393	-596.381	1.580
	Qt	0.963	0.047	0.391	-595.099	2.861
	Pf	0.928	0.047	0.394	-597.961	0.000
DW/AFDW	Ln	0.999	0.005	0.004	-1432.479	5.734
	Qd	0.999	0.005	0.004	-1434.282	3.932
	Cub	0.999	0.005	0.004	-1438.214	0.000
	Qt	0.999	0.005	0.004	-1436.168	2.045
	Pf	0.999	0.005	0.004	-1437.069	1.145

Figure 1

Map of the Caraguatatuba Bay, southeastern Brazil. Sampling areas are highlighted in grey.

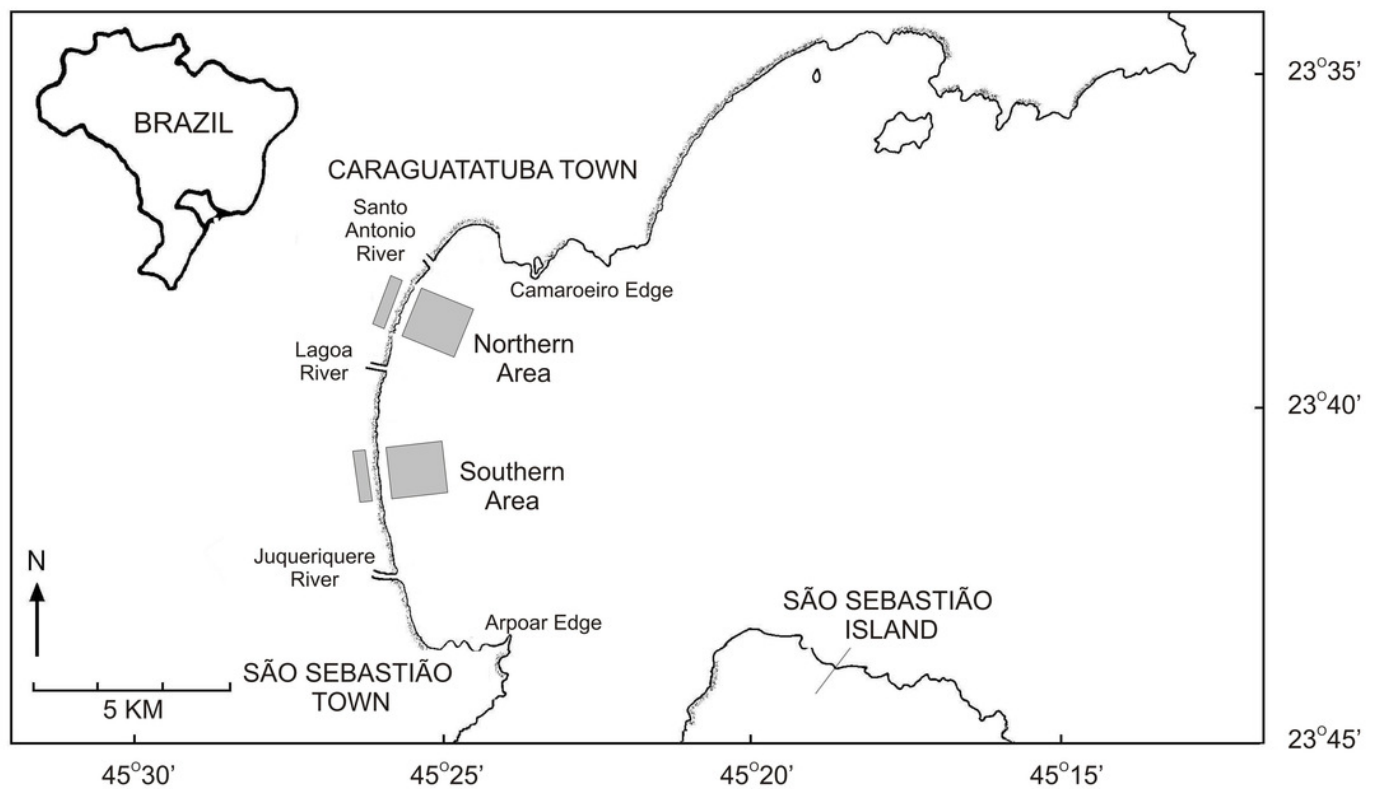


Figure 2

Tivela mactroides. Scheme indicating the measurements taken from the shells (shell length, L; height, H; and width, W). (A) frontal view. (B) lateral view.

Tivela mactroides

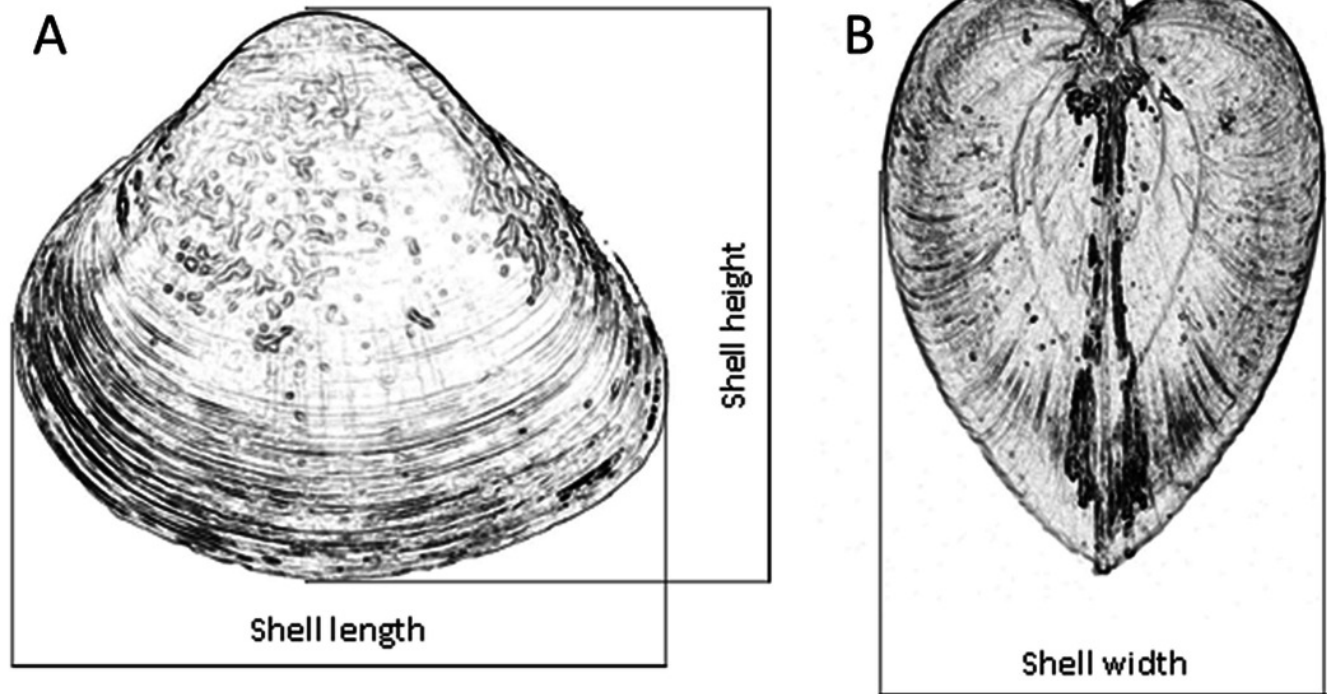


Figure 3

Tivela mactroides. Graphical representations of the allometric relationships.

Continuous lines represent the most suitable model describing the morphometric relationship. N: number of individuals analyzed. SE: standard error. AFDW: ash-free dry weight. (A) shell width ~ shell length. (B) shell height ~ shell length. (C) shell weight ~ shell length. (D) dry weight ~ shell length. (E) ash free dry weight ~ shell length. (F) shell height ~ shell width. (G) dry weight ~ shell weight. (H) ash free dry weight ~ dry weight.

