

Supplementary Materials for  
**Dental form and function in the early feeding diversification of dinosaurs**

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**This PDF file includes:**

Supplementary Text  
Figs. S1 to S12  
Tables S1 to S21  
References

## Supplementary Text

### Dinosaur tooth 3D modelling

Eleven early branching members of Dinosauria were included in this study. Maxillary or dentary teeth were selected to build the models based on preservation. Premaxillary teeth were omitted as their morphology differs from that of the cheek dentition in most taxa. Models were created from skull CT data segmented in Avizo Lite 9.5 (Thermo Fisher Scientific) and tooth surface models were cleaned and edited in Blender 2.92 (Blender Foundation) to recreate dental morphology guided by relevant literature. None of the dinosaur species studied show notable dental wear, except for *Manidens* and, to a lesser extent, *Lesothosaurus*.

*Manidens condorensis* is a heterodontosaurid ornithischian from the Early Jurassic of South America (48, 49). The dentition of *Manidens* shows marked heterodonty, with diamond-shaped maxillary teeth and closely packed hand-shaped dentary teeth, as well as a pair of anterior dentary caniniform teeth (41, 48, 65). Our reconstruction of *Manidens* tooth shape is based on CT scans of the holotype MPEF-PV 3211 (49), from which maniform dentary teeth were selected due to their autapomorphic and complex morphology. In particular, the right ninth dentary tooth was segmented, which lacks any wear facets (49), and later repaired based on the rest of the dentary dentition from the CT data as well as images and descriptions of *Manidens* dentary teeth (41, 48).

*Lesothosaurus diagnosticus* is an early branching ornithischian from the Early Jurassic of southern Africa (15, 87, 88). The teeth of *Lesothosaurus* have triangular crowns with multiple cusps along the mesiodistal margin. The crown is constricted at the base and shows the characteristic cingulum present in many ornithischians. *Lesothosaurus* does not show evident heterodonty along the tooth row (61). The tooth model corresponding to this species was obtained from the left dentary of specimen BP/1/7853 (61). The fourteenth dentary tooth was segmented for being the best preserved and lacking any notable wear (61). Minimal repair was guided by other teeth in BP/1/7853 and other specimens (15, 61).

*Eodromaeus murphi* is an early branching saurischian or theropod from the Carnian (Late Triassic) of South America (8, 13). *Eodromaeus* is characterized by having curved and laterally compressed teeth which are finely serrated on their mesial and distal carinae. Dental morphology does not vary along the toothrow, although some maxillary teeth are enlarged into caniniforms (13). The *Eodromaeus* tooth model was obtained from CT scans of the holotype PVSJ 560, represented by disarticulated skull material. The best-preserved dentary tooth, belonging to the right mandible, was segmented and the resulting surface file was cleaned and repaired based on descriptions and images of the dentition (13).

*Herrerasaurus ischigualastensis* is a herrerasaurid dinosaur from the Carnian (Late Triassic) of South America (12, 89). Its teeth are laterally compressed and curved, with a rounded mesial surface on the basal half of the crown and a sharp distal carina. Fine serrations are present all along the distal carina and along the apical part of the mesial carina. Dental morphology is relatively uniform along the toothrow, although size variation is notable among the maxillary dentition (12). A CT dataset of the specimen PVSJ 407 was the basis of our *Herrerasaurus* tooth reconstruction. The eleventh right dentary tooth was segmented due to its better preservation and later edited to remove resolution artefacts and add morphological details described in the original study of the species (12).

*Buriolestes shultzi* is the earliest branching sauropodomorph from the Carnian (Late Triassic) of South America (11, 90). The dentition of *Buriolestes* is characterized by the presence of curved, laterally compressed teeth bearing fine serrations along the distal and mesial carinae. Both maxillary and dentary toothrows show size heterodonty (11, 90). The *Buriolestes* tooth model was created from CT data of CAPP/UFMS 0035 (56), selecting a middle right maxillary tooth for its preservation. The segmented surface file was cleaned and mirrored to resemble a right dentary tooth. It was also modified to reflect the characteristic set of traits of the species based on descriptions and photographs (11, 90).

*Eoraptor lunensis* is an early dinosaur from the Carnian (Late Triassic) of South America (19), usually recovered as one of the earliest branching sauropodomorphs (8, 11, 90, 91) but occasionally classified as an early theropod (7). The dentition of *Eoraptor* is composed of laterally compressed tooth crowns that show a basal constriction and are only slightly recurved. Tooth crowns show a labial rounded eminence and serrations along the distal carina and on the apical half of the mesial carina (19). CT scans of the skull of the holotype PVSJ 512 were segmented to isolate a representative tooth model, choosing the well-preserved sixth left maxillary tooth. The model was edited to include fine details like the serrations, which are not captured in the CT scans, following photographs and descriptions of the specimen (19).

*Saturnalia tupiniquim* is an early member of Sauropodomorpha from the Carnian (Late Triassic) of South America (92). Several aspects of the dentition of *Saturnalia* are unknown due to the generally poor preservation of the cranial material. However, well-preserved maxillary and dentary crowns show a basal constriction followed by a mesiodistal expansion of the crown, which tapers apically. The mesial margin is anteriorly convex while the distal margin is straight to sigmoid, producing a slight posterior curvature of the teeth. The distal margins of the best-preserved teeth show fine denticles (20). The *Saturnalia* tooth model was obtained from CT data of the specimen MCP-3845-PV, selecting a well-preserved right dentary tooth, approximately the ninth along the tooth row. The surface model was cleaned, and details missed from the CT scans were modelled according to published descriptions and images (20, 57).

*Thecodontosaurus antiquus* is an early branching sauropodomorph from the Rhaetian (Late Triassic) of Europe (40, 93). The dentition of *Thecodontosaurus* is composed of lanceolate crowns with coarse denticles on the mesial and distal margins. The maxillary and dentary crowns are slightly posteriorly inclined and show a basal constriction and a tapering apex. Both the labial and lingual surfaces bear a central apicobasal eminence which is more developed on the labial side (40, 93). The *Thecodontosaurus* tooth model was created from CT scans of the specimen BRSUG 28221, a fragmentary left maxilla (40). A posterior left maxillary tooth was segmented and later cleaned to remove segmentation artefacts.

*Plateosaurus engelhardti* is an early sauropodomorph dinosaur from the Norian (Late Triassic) of Europe (38). *Plateosaurus* maxillary and dentary teeth are lanceolate, with convex mesial and distal margins in labial view. These margins bear coarse and prominent denticles along their length, as well as a labial eminence along the apicobasal axis of the crown. Tooth crowns are almost straight and constricted at their base (38). The source of the *Plateosaurus* tooth model is CT data of the specimen MB.R.1937, a complete skull (47), from which the fourteenth left dentary tooth was isolated. The surface modelled was mirrored to the right counterpart and was cleaned and smoothed to remove resolution artefacts. Fine dental details were modelled based on photographs, drawings and descriptions of *P. engelhardti* specimens (14, 38) and other *Plateosaurus* species (94, 95).

*Massospondylus carinatus* is a massospondylid sauropodomorph from the Early Jurassic of southern Africa (37). The cheek teeth of *Massospondylus* are lanceolate, apicobasally elongate and constricted at the base. Crowns bear apically oriented denticles on the apical part of the mesial and distal carinae. Variation in size is seen along the tooth row as crown decrease in height towards the posterior end (37, 96). Our *Massospondylus* tooth reconstruction is based on CT data of the specimen BP/1/5241, a complete cranium (96). The eighth left maxillary tooth was segmented due to its preservation. The surface model was smoothed and the denticles, not captured in the scans, were modelled based on available information of *Massospondylus* specimens (37, 96).

*Ngwevu intloko* is a massospondylid sauropodomorph from the Early Jurassic of southern Africa (39, 96). Dental morphology in *Ngwevu* is similar to that of *Massospondylus* in having maxillary and dentary teeth with lanceolate and elongated crowns that are constricted at their base. Coarse denticles are present on the apical part of the crowns. The *Ngwevu* tooth model was obtained from CT scans of the skull of the holotype BP/1/4779. The eleventh right maxillary tooth was segmented and later cleaned, based on photographs of the specimen (39), and mirrored to resemble a right dentary crown.

### Constraint test

A test was performed to assess the impact of varying the number of constraint nodes and the position of the reference point in the multi-point constraint at the base of the crown. The *Iguana iguana* tooth model was tetrameshed in Hypermesh 2017 (Altair), resulting in a 3D mesh composed of 3296634 C3D4 tetrahedral elements. Material properties of bovine dentine – Young’s modulus of 21 GPa and a 0.31 Poisson’s ratio (68) – were assigned to the model. Boundary conditions and loads were applied were applied in Abaqus 6.14 (Simulia). We used a multi-point constraint to ‘fix’ the base of the tooth crown during the simulation. This involves a defined number of constraint nodes around the tooth base connected to a fixed reference point node. The test looked at the effect on tooth stress distribution and magnitude of using different numbers of nodes to constrain movement at the base of the tooth crown. A vertical and basally-oriented force of 100 N was applied to a node at the tip of the tooth crown. Ten simulations were performed, varying the number of constraint nodes: 20, 40, 60, 80 and all (378) nodes – and the position of the reference point along the apicobasal (y) axis from either immediately below the crown base surface to a distance equal to the crown height, representing the root. The multi-point constraint was constrained in all degrees of freedom, both in translation and rotation. Von Mises stress and element volume were obtained from the analyses, and these were used to calculate the mesh-weighted arithmetic mean Von Mises stress of models (30).

The results of the constraint test confirm that varying the position of the constraint reference point has no effect on stress values, and variations in this magnitude resulting from increasing the number of constraint nodes is not statistically significant, with error percentages below 1% (fig. S1, table S3). This demonstrates that the number of slave nodes and the position of the reference point in the multi-point constraint has no significant impact on stress magnitude and distribution in our FE models.

### Convergence test

A convergence test was carried out to decide the element size and approximate number of elements of the mesh that should be used in the simulations, in order to minimize errors due to mesh resolution across models. The *Iguana iguana* tooth model was used in the test and the 2D

mesh was remeshed in Blender 2.92 with varying edge lengths to generate seven surface models with different resolution (table S4). These surface models were tetrameshed in Hypermesh 2017, generating meshes composed of different number of C3D4 elements (table S4), and material properties of bovine dentine were assigned to them (68). Boundary conditions and loads were applied on Abaqus 6.14, using a multipoint constraint composed of 20 nodes located around the exterior edge of the crown base and a reference point placed right below the centre of the base. The base was constrained in all degrees of freedom and a vertical, basally oriented force of 100 N was applied on a node at the tip of the crown. Von Mises stress and element volume was requested as analysis output parameters to calculate the mesh-weighted arithmetic mean of Von Mises stress (30).

Meshed ranging from approximately 10 million to 600,000 elements, generated with element edge lengths from 0.05 to 0.2 mm, yield similar von Mises stress values (errors below 3%, fig. S2, table S4), indicating that convergence is achieved within this range of mesh resolution.

### Phylogeny time-calibration

We built an informal supertree combining the phylogenetic topologies of extant squamates (97), crocodylians (98) and non-avian dinosaurs (9) (fig. S11). We time calibrated the phylogeny using occurrence data from the Paleobiology Database (fossilworks.org) and original sources (13, 99, 100), constraining the calibration with minimum node ages of eight clades (98, 101, 102) (table S5). Time calibration was performed using the timePaleoPhy function of the paleotree package (103) in R.

### Machine learning algorithms information

The nine machine learning algorithms used in this study are briefly described below based on Kuhn and Johnson (85).

Classification and Regression Trees (CART) partition the data into smaller groups and fit a simple model to each of them (104). These subgroups are more homogeneous in the response variable. CART are graphically represented with decision trees. The parameters that can be tuned are the complexity of the tree (cost complexity), the minimum number of splits and the maximum depth of the tree.

Linear Discriminant Analysis (LDA) finds the linear combination of predictor variables that maximizes between-group variance in relation to within-group variance (105, 106). Thus, LDA looks for the combination of predictors that maximizes the separation between the centre of the groups and minimizes the variation within each group. LDA does not require tuning.

Mixture Discriminant Analysis (MDA) is an extension of LDA that allows a set number of subclasses within each class (107). The tuning parameter is the number of subclasses.

Support Vector Machines (SVM) separate classes of the dependent variable by defining boundaries between groups and maximizing the margin (i.e., the distance between the boundary and the closest observation within the training sample) (108). Boundaries are drawn using kernel functions which can be linear, polynomial or radial. The main tuning parameter is the cost, which is the penalty for misclassification caused by reducing the margin. Here, we use linear (SVML) and radial (SVMR) support vector machines.

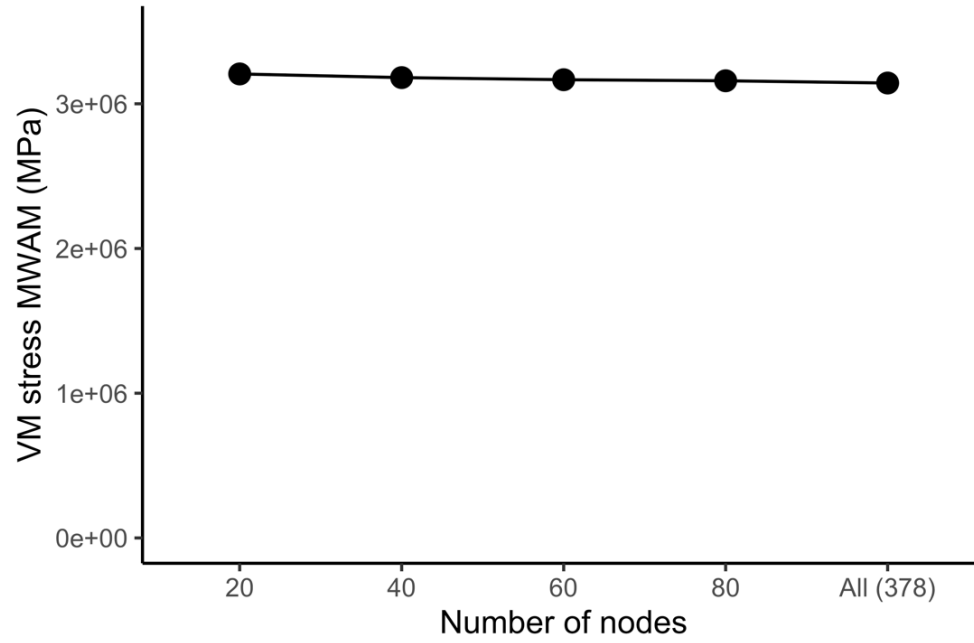
k-Nearest Neighbor (kNN) establishes classifications based on the distance of a sample to the neighbors of each class (109). The sample is classified in the group that is more frequent

among the neighbors within a given distance. The tuning parameter for this algorithm is the number of neighbors ( $k$ ).

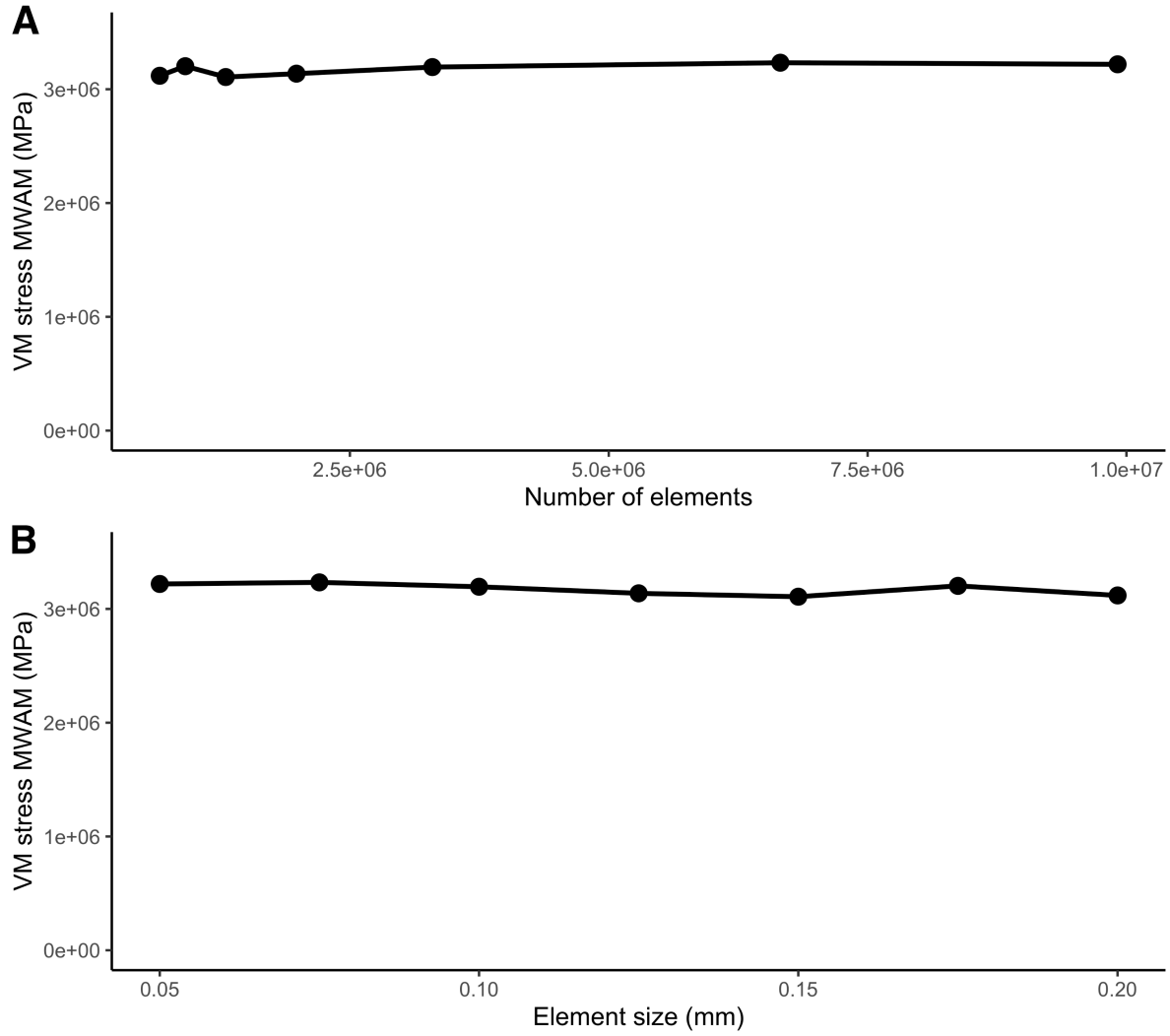
Naïve Bayes (NB) applies Bayes' probability rule to calculate the probability of belonging to a class given the predictor variables in the dataset (110). The model assumes that the predictor variables are independent. The tuning parameters for NB deal with the use of kernel density estimates and Laplace correction, that prevents predictors without train set samples to make the posterior probability zero.

Random Forest (RF) creates a series of classification trees that are obtained from bootstrap resamples of the data, making the trees independent and decorrelated. The class probability of a sample is derived from the distribution of individual classifications made by each tree in the forest (111). The tuning parameter is the number of randomly selected predictor variables to choose from at each split ( $m_{try}$ ).

Neural Networks (NNet) combine predictor variables into a set of hidden units that conform one or more hidden layers (112). Information from the hidden layer(s) is combined to produce the classification output. A cost function is used to assess classification error and improve the model via back propagation. The main tuning parameters are the number of hidden units (size) and the weight decay (decay), which penalizes overfitting.

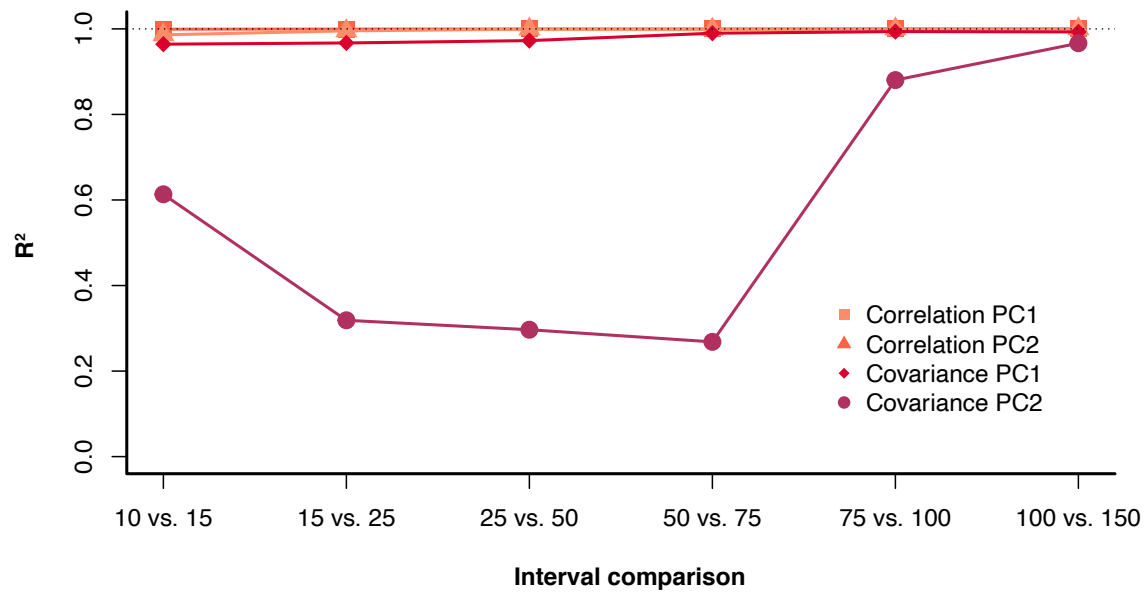


**Fig. S1. Constraint test.** Variation in von Mises stress mesh-weighted arithmetic mean (MPa) with varying number of constrained nodes around the base of the tooth FE model of *Iguana iguana*.

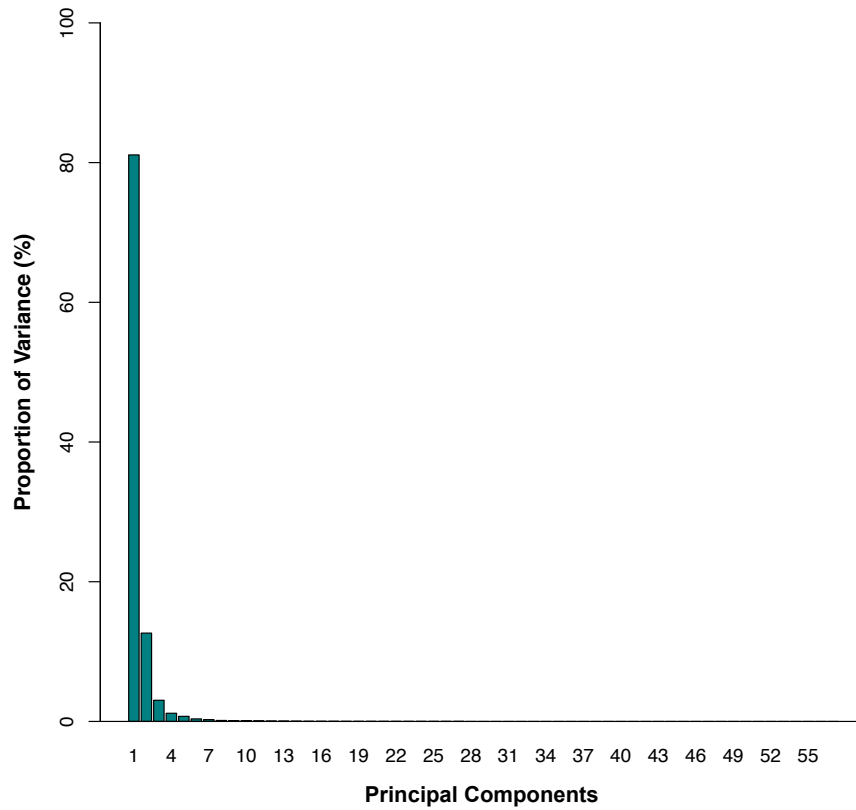


**Fig. S2. Convergence test.** Variation in von Mises stress mesh-weighted arithmetic mean (MPa) with varying a number of tetrahedral elements and b corresponding mean element size in mm in the tooth FE model of *Iguana iguana*.

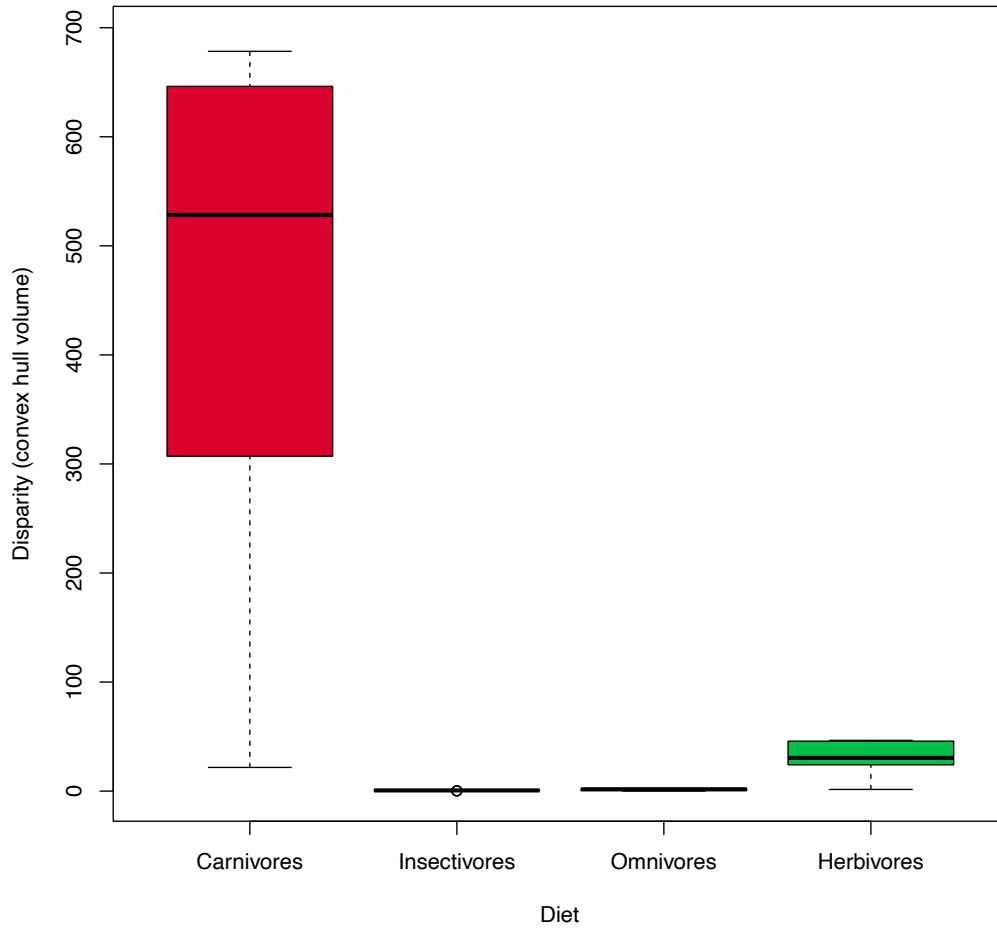




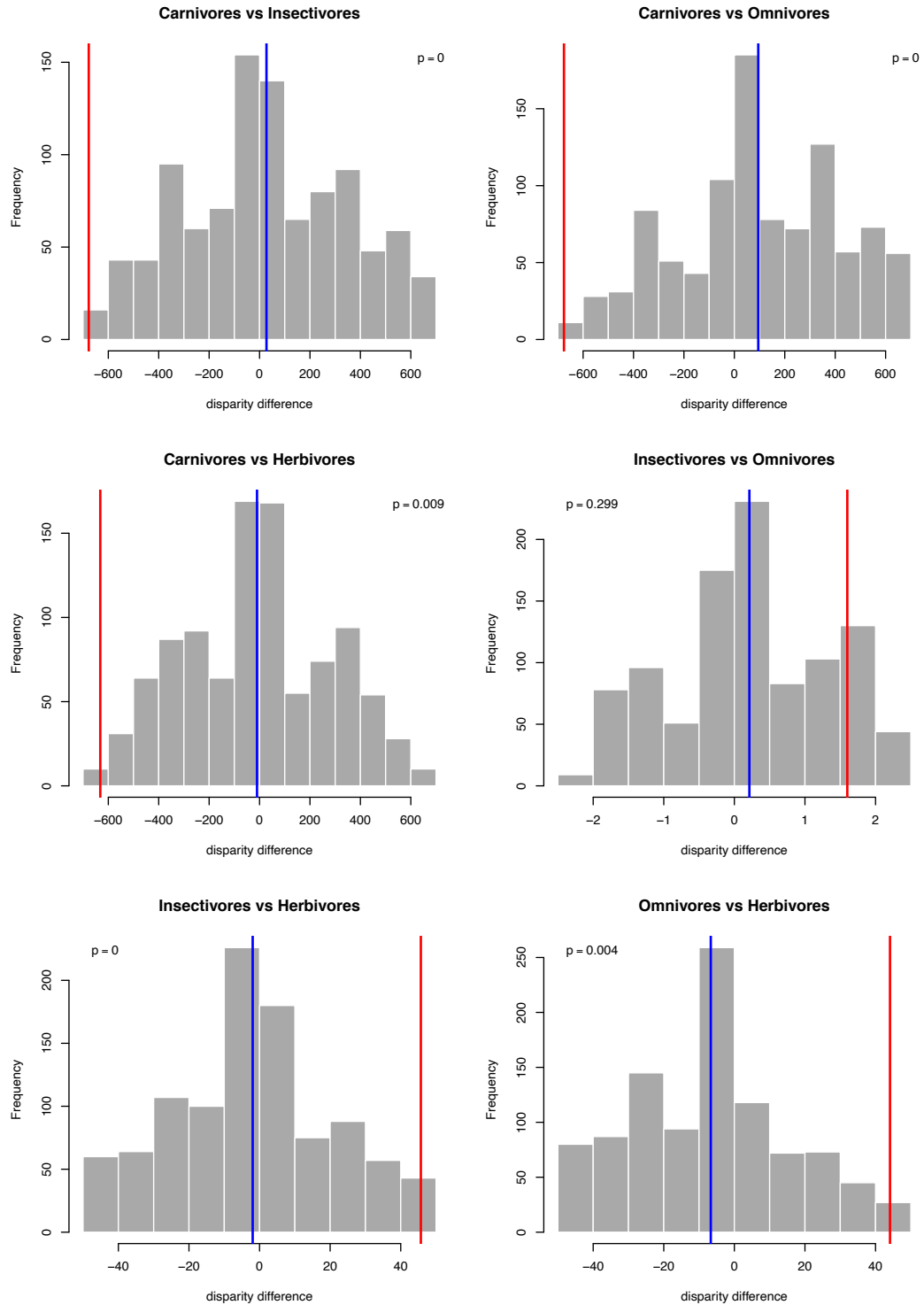
**Fig. S3. Convergence of intervals.**  $R^2$  values of pairwise comparisons of PCAs using different number of intervals and either correlation or covariance matrices.



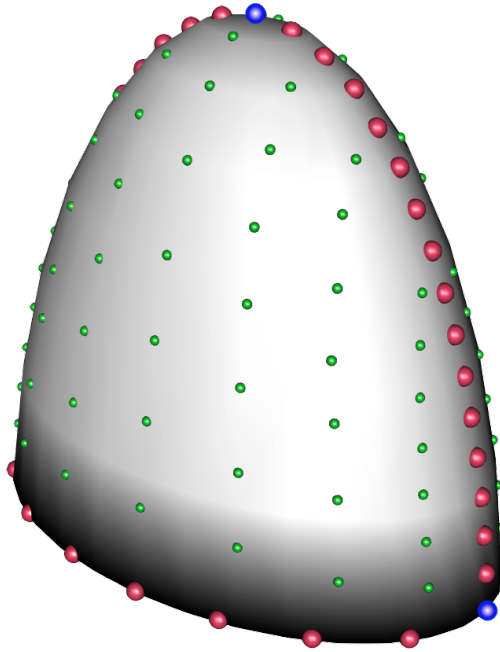
**Fig. S4. Proportion of variance explained by each Principal Component resulting from the PCA on the stress intervals matrix.**



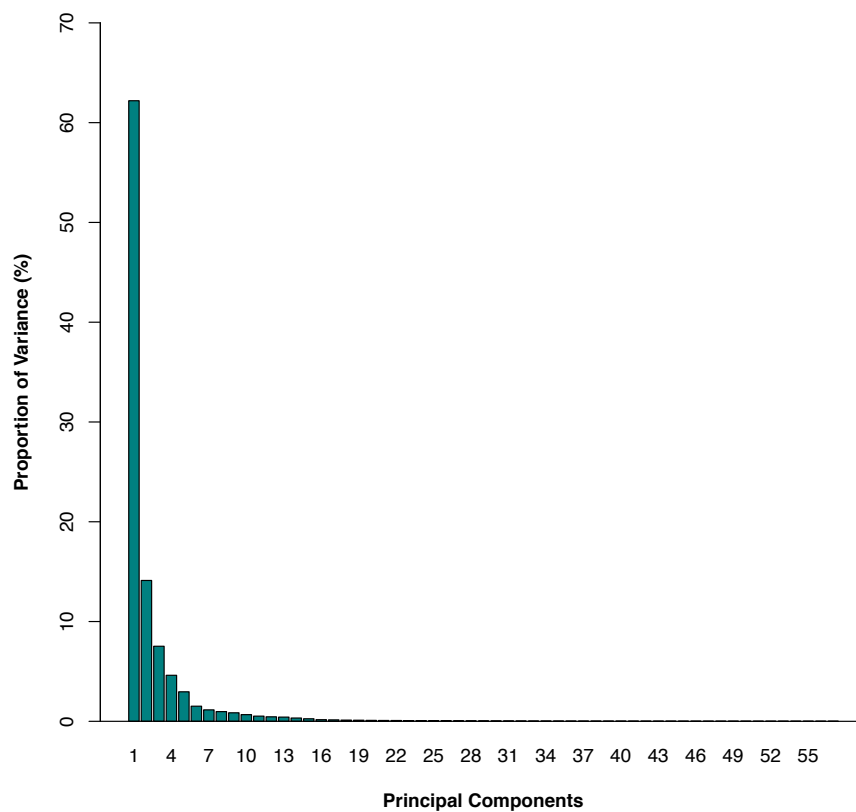
**Fig. S5. Biomechanical disparity per dietary group.** Disparity is measured as the convex hull volume of each group considering the first two PC axes of the biomechanical space. Disparity for durophages could not be calculated due to their small sample size.



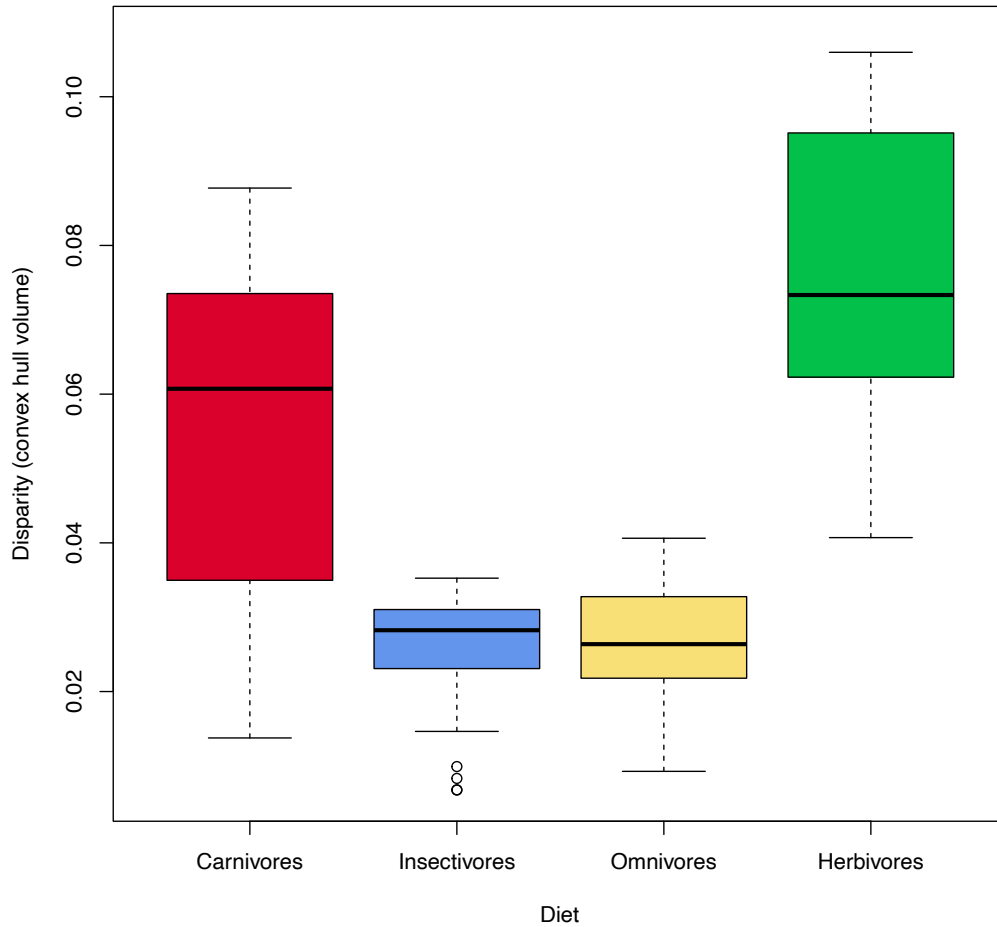
**Fig. S6. Permutation test comparisons of biomechanical disparity between dietary groups.** Lines show observed (red) and expected (blue) disparity differences and p-values (top of the histograms) indicate if differences are statistically significant when  $p < 0.05$ .



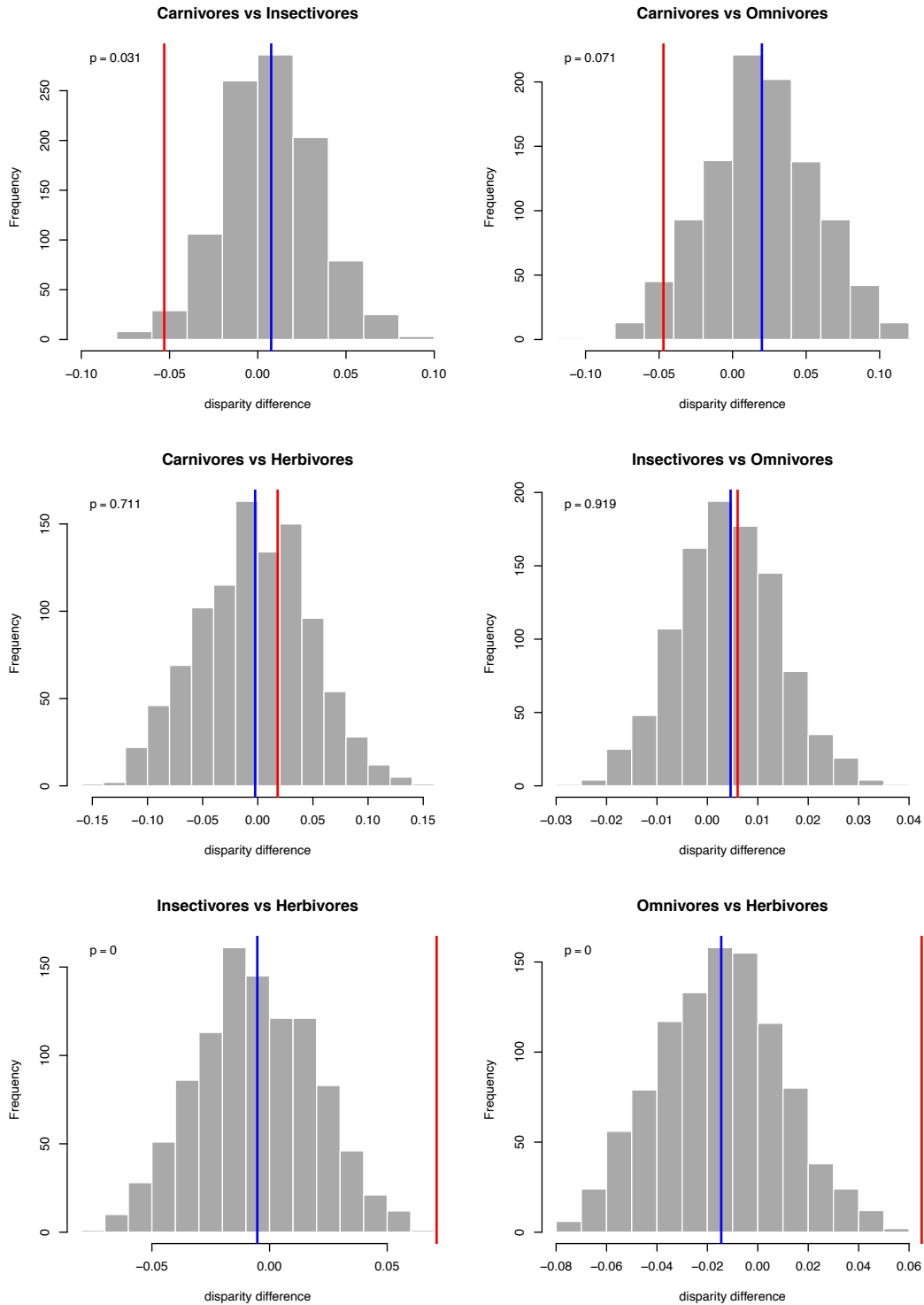
**Fig. S7. Template used to transfer surface semi-landmarks to tooth models in mesiolabial view.** Fixed landmarks are shown in blue, curve semi-landmarks in red and surface semi-landmarks in green.



**Fig. S8. Proportion of variance explained by each Principal Component resulting from the PCA on the Procrustes alignment.**

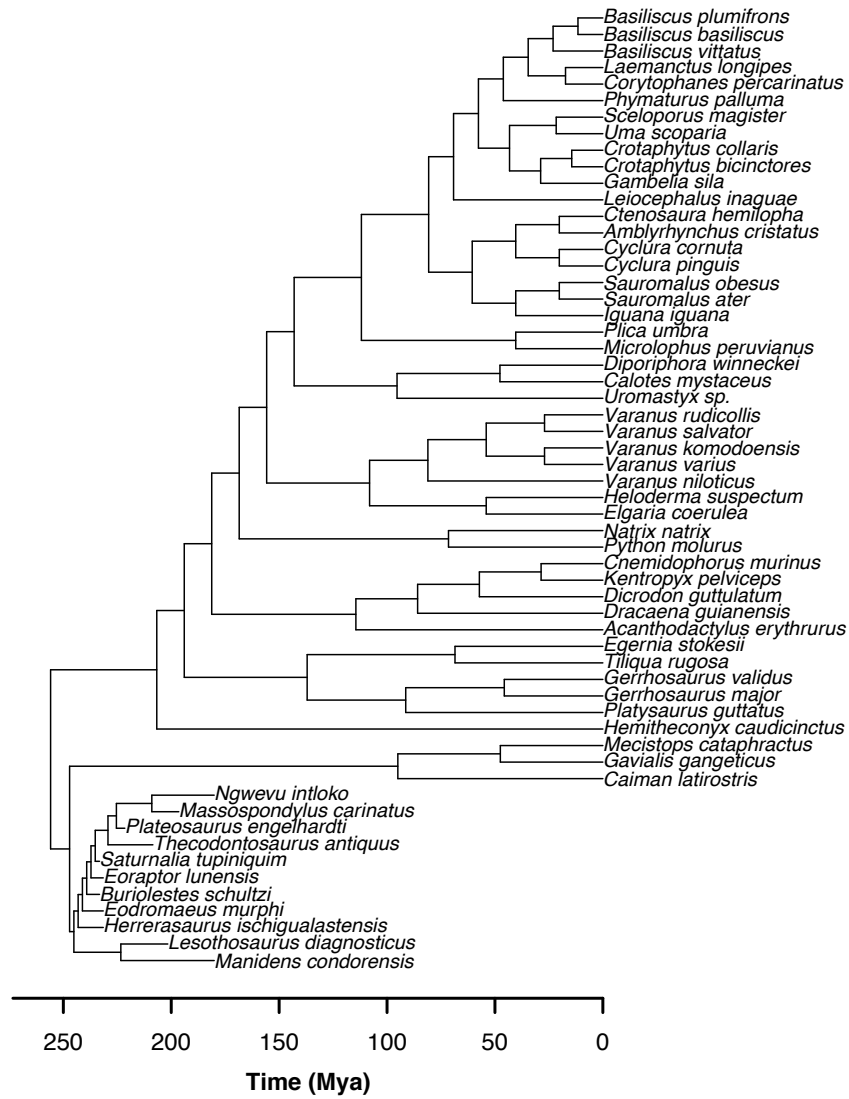


**Fig. S9. Morphological disparity per dietary group.** Disparity is measured as the convex hull volume of each group considering the first two PC axes of the morphological space. Disparity for durophages could not be calculated due to their small sample size.

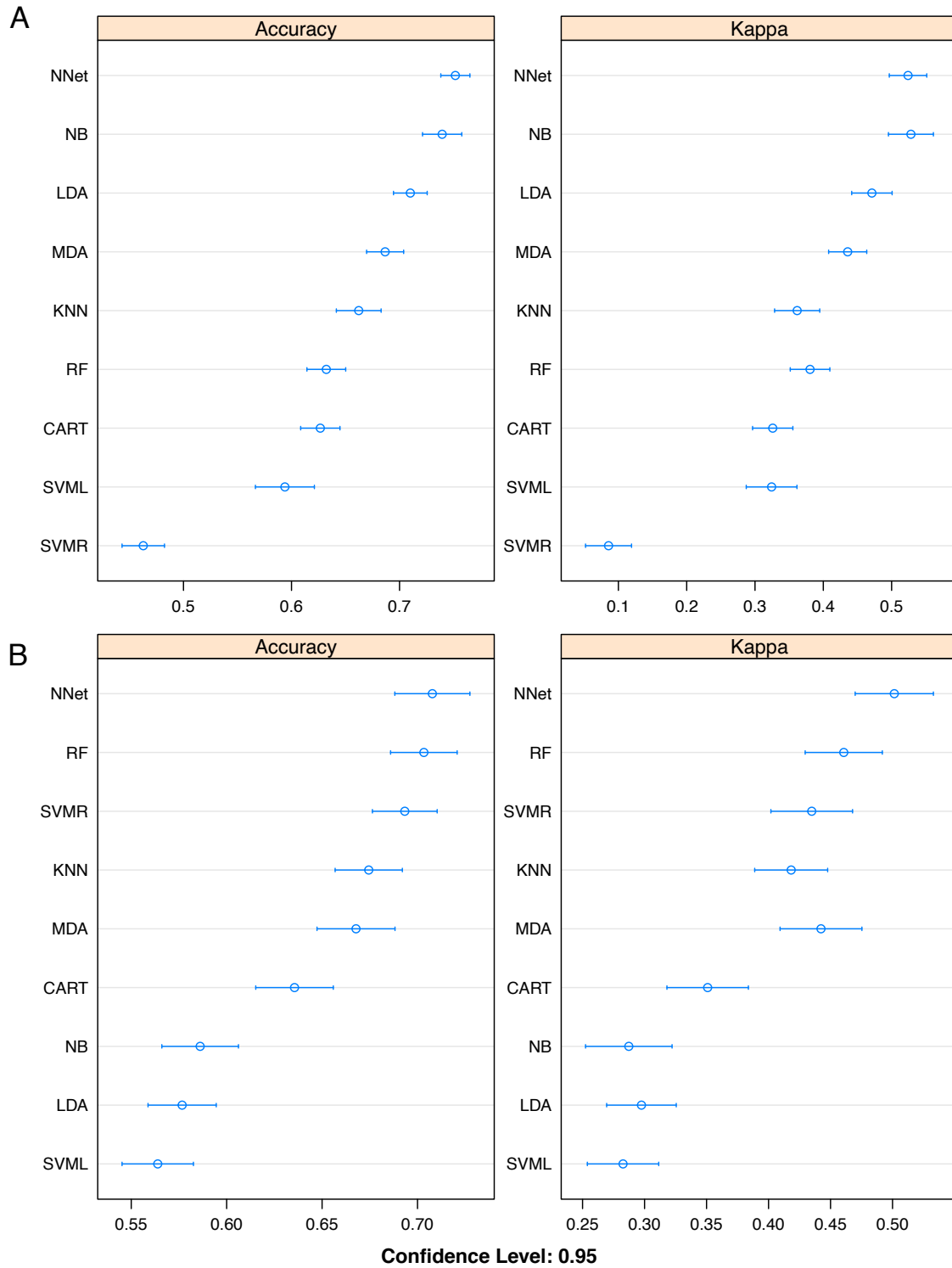


**Fig. S10. Permutation test comparisons of morphological disparity between dietary groups.** Lines show observed (red) and expected (blue) disparity differences and p-values (top left of the histograms) indicate if differences are statistically significant when  $p < 0.05$ .





**Fig. S11. Time-calibrated phylogeny of Sauropsida with the 58 taxa included in the study.**



**Fig. S12. Accuracy and Kohen’s Kappa comparisons of nine machine learning algorithms tested on the (A) biomechanical and (B) morphological data.**

**Table S1. Specimen and data source information.** Dietary information compiled from published quantitative datasets and ecological studies of lizards (113, 114), snakes (115, 116) and crocodylians (117–119). Predators were split into carnivores, insectivores and durophages depending on whether they preferentially prey on vertebrates, arthropods or hard-shelled animals, respectively (23).

Species	Specimen	Tooth position	Clade	Data type	Diet	Source	Author	Ref.
<i>Crotaphytus bicinctores</i>	CAS 200859	Right Dentary 19	Squamata	Surface model	Carnivore	MorphoSource	Keegan Melstrom	(23)
<i>Gavialis gangeticus</i>	UF H 118998	Left Dentary 10	Crocodylia	CT data	Carnivore	MorphoSource	David Blackburn	
<i>Heloderma suspectum</i>	CAS 159492	Right Dentary 4	Squamata	Surface model	Carnivore	MorphoSource	Keegan Melstrom	(23)
<i>Mecistops cataphractus</i>	TMM M-3529	Right Dentary 11	Crocodylia	CT data	Carnivore	MorphoSource	Jessica Maisano	
<i>Natrix natrix</i>	UMMZ H 65465	Right Dentary 9	Squamata	CT data	Carnivore	MorphoSource	Gregory Schneider	
<i>Python molurus</i>	UF H 190353	Right Dentary 14	Squamata	CT data	Carnivore	MorphoSource	David Blackburn	
<i>Varanus komodoensis</i>	TNHC 95803	Right Dentary 8	Squamata	CT data	Carnivore	MorphoSource	Jessica Maisano	
<i>Varanus rudicollis</i>	UCMP 137816	Right Dentary 10	Squamata	Surface model	Carnivore	MorphoSource	Keegan Melstrom	(23)
<i>Varanus salvator</i>	PIMUZ 7825	Right Dentary 3	Squamata	Surface model	Carnivore	MorphoSource	Eva Herbst	(120)
<i>Varanus varius</i>	MVZ 77092	Right Dentary 10	Squamata	Surface model	Carnivore	MorphoSource	Keegan Melstrom	(23)
<i>Caiman latirostris</i>	UMMZ H 155287	Left Dentary 18	Crocodylia	CT data	Durophage	MorphoSource	Gregory Schneider	
<i>Dracaena guianensis</i>	MVZ 79247	Right Dentary 9	Squamata	Surface model	Durophage	MorphoSource	Keegan Melstrom	(23)
<i>Varanus niloticus</i>	MVZ 68534	Right Dentary 9	Squamata	Surface model	Durophage	MorphoSource	Keegan Melstrom	(23)
<i>Amblyrhynchus cristatus</i>	MVZ 67721	Right Dentary 8	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Cyclura cornuta</i>	UCMP 123055	Right Dentary 19	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Dicrodon guttulatum</i>	MVZ 85400	Left Dentary 15	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Egernia stokesii</i>	MCZ R-33105	Right Dentary 14	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Iguana iguana</i>	UMNH 8084	Right Dentary 13	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Phymaturus palluma</i>	MVZ 137647	Right Dentary 13	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Sauromalus ater</i>	MVZ 100404	Right Dentary 13	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Sauromalus obesus</i>	UCMP 137811	Right Dentary 17	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Tiliqua rugosa</i>	MCZ R-24456	Right Dentary 10	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Uromastix sp.</i>	UCMP 118912	Right Dentary 10	Squamata	Surface model	Herbivore	MorphoSource	Keegan Melstrom	(23)
<i>Basiliscus vittatus</i>	UCMP 137748	Right Dentary 18	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Calotes mystaceus</i>	CAS 243761	Right Dentary 14	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Corytophanes percarinatus</i>	UCMP 123057	Right Dentary 19	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Crotaphytus collaris</i>	UCMP 141134	Right Dentary 15	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Elgaria coerulea</i>	CAS 216644	Right Dentary 19	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Gambelia sila</i>	CAS 141318	Right Dentary 17	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)

<i>Hemitheconyx caudicinctus</i>	CAS uncatalogued	Right Dentary 23	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Kentropyx pelviceps</i>	MVZ 174866	Right Dentary 18	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Laemanctus longipes</i>	UCMP 129880	Right Dentary 18	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Plica umbra</i>	CAS 93242	Right Dentary 18	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Sceloporus magister</i>	CAS 200862	Right Dentary 20	Squamata	Surface model	Insectivore	MorphoSource	Keegan Melstrom	(23)
<i>Acanthodactylus erythrurus</i>	SDNHM 65156	Right Dentary 18	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Basiliscus basiliscus</i>	MVZ 79579	Right Dentary 14	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Basiliscus plumifrons</i>	MCZ R-19487	Right Dentary 17	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Cnemidophorus murinus</i>	OU 39632	Right Dentary 10	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Ctenosaura hemilopha</i>	CAS 46399	Right Dentary 13	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Cyclura pinguis</i>	KU 272292	Left Dentary 22	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Diporiphora winneckeii</i>	MCZ R-35222	Right Dentary 9	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Gerrhosaurus major</i>	UCMP 137878	Right Dentary 16	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Gerrhosaurus validus</i>	MCZ R-50973	Right Dentary 21	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Leiocephalus inaguae</i>	MCZ R-154263	Right Dentary 20	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Microlophus peruvianus</i>	UCMP 141136	Right Dentary 17	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Platysaurus guttatus</i>	MCZ R-44415	Right Dentary 16	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Uma scoparia</i>	CAS 42076	Right Dentary 24	Squamata	Surface model	Omnivore	MorphoSource	Keegan Melstrom	(23)
<i>Manidens condorensis</i>	MPEF-PV 3211	Right Dentary 9	Dinosauria	CT data	Unknown	From author	Marcos Becerra	(49)
<i>Lesothosaurus diagnosticus</i>	BP/1/7853	Left Dentary 14	Dinosauria	CT data	Unknown	From author	Lara Sciscio	(61)
<i>Eodromaeus murphi</i>	PVSJ 560	Right Dentary ?	Dinosauria	CT data	Unknown	From author	Paul Barrett, David Button, Ricardo Martínez, Laura Porro	
<i>Herrerasaurus ischigualastensis</i>	PVSJ 407	Right Dentary 11	Dinosauria	CT data	Unknown	From author	Paul Barrett, David Button, Ricardo Martínez, Laura Porro	
<i>Buriolestes schultzi</i>	CAPPA/UFFS M 0035	Right Maxillary 8	Dinosauria	CT data	Unknown	From author	Rodrigo Müller	(56)
<i>Eoraptor lunensis</i>	PVSJ 512	Left Maxillary 6	Dinosauria	CT data	Unknown	Digimorph	Jessica Maisano	
<i>Saturnalia tupiniquim</i>	MCP-3845-PV	Right Dentary 9?	Dinosauria	CT data	Unknown	From author	Mario Bronzati	(20)
<i>Thecodontosaurus antiquus</i>	BRSUG 28221	Left Maxillary ?	Dinosauria	CT data	Unknown	From author	Antonio Ballell	(40)
<i>Plateosaurus engelhardti</i>	MB.R.1937	Left Dentary 14	Dinosauria	CT data	Unknown	From author	Lawrence Witmer	(47)
<i>Massospondylus carinatus</i>	BP/1/5241	Left Maxillary 8	Dinosauria	CT data	Unknown	From author	Kimberley Chapelle	(96)
<i>Ngwevu intloko</i>	BP/1/4779	Right Maxillary 11	Dinosauria	CT data	Unknown	From author	Kimberley Chapelle	(39)

**Table S2. Details of 2D, 3D meshes and original surface area and volume of tooth models.**

Model	2D mesh	3D mesh		Tooth surface area (mm <sup>2</sup> )	Tooth volume (mm <sup>3</sup> )
	No. triangles	No. elements	No. nodes		
<i>Acanthodactylus erythrurus</i>	121610	1443842	212665	8.205E-05	5.373E-08
<i>Amblyrhynchus cristatus</i>	125012	1273969	183360	3.656E-04	3.914E-07
<i>Basiliscus basiliscus</i>	124112	1369388	199576	4.260E-04	5.851E-07
<i>Basiliscus plumifrons</i>	122612	1347262	196160	4.735E-04	7.052E-07
<i>Basiliscus vittatus</i>	119980	1292079	187555	1.815E-04	1.614E-07
<i>Buriolestes schultzi</i>	122458	1334852	194036	3.883E-04	5.038E-07
<i>Caiman latirostris</i>	122950	1367401	199419	8.243E-03	5.577E-05
<i>Calotes mystaceus</i>	125158	1444632	274444	1.373E-04	9.390E-07
<i>Cnemidophorus murinus</i>	120956	1484242	219617	9.063E-05	6.382E-08
<i>Corytophanes percarinatus</i>	120138	1350974	197456	3.779E-04	4.727E-07
<i>Crotaphytus bicinctores</i>	122808	1492185	220402	2.029E-04	1.995E-07
<i>Crotaphytus collaris</i>	122264	1451999	213902	3.864E-04	5.283E-07
<i>Ctenosaura hemilopha</i>	122524	1342753	195501	7.170E-05	4.026E-08
<i>Cyclura cornuta</i>	122938	1345600	195879	4.255E-04	5.398E-07
<i>Cyclura pinguis</i>	121548	1306602	189644	2.554E-04	2.698E-07
<i>Dicrodon guttulatum</i>	121800	1461539	215654	1.620E-04	1.449E-07
<i>Diporiphora winneckeii</i>	121556	1438749	211761	1.490E-04	1.197E-07
<i>Dracaena guianensis</i>	117266	1481481	220155	2.034E-02	2.023E-04
<i>Egernia stokesii</i>	121760	1420876	208723	5.371E-05	2.986E-08
<i>Elgaria coerulea</i>	123124	1401739	205190	2.038E-04	1.944E-07
<i>Eodromaeus murphi</i>	118936	1233313	177841	1.449E-03	3.090E-06
<i>Eoraptor lunensis</i>	120432	1284714	186275	3.712E-03	1.360E-05
<i>Gambelia sila</i>	120162	1391578	204212	3.219E-04	3.915E-07
<i>Gavialis gangeticus</i>	123510	1255332	180524	2.197E-02	1.784E-04
<i>Gerrhosaurus major</i>	117818	1376988	202353	6.130E-04	1.170E-06
<i>Gerrhosaurus validus</i>	121086	1418604	208485	9.711E-05	7.072E-08
<i>Heloderma suspectum</i>	122088	1217125	174527	2.273E-03	5.403E-06
<i>Hemitheconyx caudicinctus</i>	121620	1417703	208288	4.086E-05	1.978E-08
<i>Herrerasaurus ischigualastensis</i>	119614	1259931	182248	1.116E-02	6.681E-05
<i>Iguana iguana</i>	122124	1300704	188477	5.405E-04	7.551E-07
<i>Kentropyx pelviceps</i>	119674	1340582	195802	5.030E-04	7.645E-07
<i>Laemantcus longipes</i>	118226	1317744	192290	3.342E-04	4.393E-07
<i>Leiocephalus inaguae</i>	121994	1429824	210213	6.427E-04	1.225E-06
<i>Lesothosaurus diagnosticus</i>	125788	1363519	198106	2.150E-03	6.799E-06
<i>Manidens condorensis</i>	120478	1200434	171777	3.123E-03	9.301E-06
<i>Massospondylus carinatus</i>	118766	1224601	176500	9.619E-03	5.640E-05
<i>Mecistops cataphractus</i>	118966	1282708	186283	1.945E-02	1.852E-04
<i>Microlophus peruvianus</i>	121658	1398018	204870	5.723E-05	2.951E-08

<i>Natrix natrix</i>	124280	1320729	190969	1.248E-04	7.861E-08
<i>Ngwevu intloko</i>	119302	1230512	177435	6.945E-03	3.465E-05
<i>Phymaturus palluma</i>	123582	1347838	196029	7.486E-05	4.280E-08
<i>Plateosaurus engelhardti</i>	121198	1266918	183008	2.011E-02	1.748E-04
<i>Platysaurus guttatus</i>	122162	1423457	209163	6.437E-05	3.841E-08
<i>Plica umbra</i>	120774	1399857	205418	1.216E-04	9.358E-08
<i>Python molurus</i>	124642	1303669	188257	1.890E-03	4.606E-06
<i>Saturnalia tupiniquim</i>	121544	1305620	189435	1.725E-03	4.765E-06
<i>Sauromalus ater</i>	123370	1356743	197672	2.237E-04	2.145E-07
<i>Sauromalus obesus</i>	124300	1410188	206282	3.109E-04	3.545E-07
<i>Sceloporus magister</i>	121002	1406836	206546	1.584E-04	1.425E-07
<i>Thecodontosaurus antiquus</i>	124202	1287487	185759	1.855E-03	4.682E-06
<i>Tiliqua rugosa</i>	122774	1530451	226861	2.144E-03	6.739E-06
<i>Uma scoparia</i>	120844	1442058	212523	6.069E-05	3.276E-08
<i>Uromastyx sp</i>	122304	1508648	223325	1.815E-04	1.548E-07
<i>Varanus komodoensis</i>	120936	1236894	178019	1.523E-02	1.046E-04
<i>Varanus niloticus</i>	121766	1413187	207495	7.054E-03	4.964E-05
<i>Varanus rudicollis</i>	121920	1273016	183866	2.640E-03	7.985E-06
<i>Varanus salvator</i>	121042	1285660	186189	5.816E-03	2.426E-05
<i>Varanus varius</i>	121722	1299564	188373	4.537E-03	1.812E-05

**Table S3. Constraint test.** Von Mises stress mesh-weighted arithmetic mean (MPa) with varying number of constrained nodes around the base of the tooth FE model of *Iguana iguana*, position of the reference point (base or root) and error between observations.

Nodes	Base constraint MWAM	Root constraint MWAM	Error %
20	3206235.68	3206235.68	0.797
40	3180873.22	3180873.22	0.462
60	3166249.23	3166249.23	0.230
80	3158976.04	3158976.04	0.496
All (378)	3143387.61	3143387.61	NA

**Table S4. Convergence test.** Von Mises stress mesh-weighted arithmetic mean (MPa) with varying number of tetrahedral elements and corresponding mean element size in mm, with error between observations.

Element size	No. elements	MWAM	Error %
0.05	9914170	3219143.48	0.437
0.075	6656992	3233270.41	1.206
0.1	3296634	3194732.59	1.842
0.125	1983843	3136949.88	0.957
0.15	1300704	3107215.41	2.975
0.175	910179	3202503.45	2.697
0.2	663545	3118399.04	NA

**Table S5. Minimum node ages of eight sauropsid clades used to constrain the time calibration of the phylogeny.**

Clade	Taxon 1	Taxon 2	Minimum age (Mya)	Ref.
Sauropsida	<i>Gavialis gangeticus</i>	<i>Basiliscus basiliscus</i>	255.9	(101)
Archosauria	<i>Gavialis gangeticus</i>	<i>Massospondylus carinatus</i>	247.1	(101)
Crocodylia	<i>Gavialis gangeticus</i>	<i>Caiman latirostris</i>	95	(98)
Squamata	<i>Hemitheconyx caudicinctus</i>	<i>Basiliscus basiliscus</i>	206.7	(101)
Scincoidea	<i>Egernia stokesii</i>	<i>Gerrhosaurus major</i>	137.02	(102)
Anguimorpha	<i>Heloderma suspectum</i>	<i>Varanus varius</i>	108.09	(102)
Iguania	<i>Uromastyx sp.</i>	<i>Basiliscus basiliscus</i>	143.05	(102)
Euiguana	<i>Iguana iguana</i>	<i>Basiliscus basiliscus</i>	80.71	(102)

**Table S6. Phylogenetic multivariate regression (PGLS) of Von Mises stress MWAM on tooth size (log tooth Centroid Size).**

	df	SS.obs	MS	F	P.val	Rsq
log.Csize	1	112859.9	112859.86	2.112862	0.3216783	0.0363579
Residual	56	2991275.9	53415.64			

**Table S7. Phylogenetic multivariate regression (PGLS) of Von Mises stress MWAM on tooth size (log tooth surface area).**

	df	SS.obs	MS	F	P.val	Rsq
log.sa	1	63387.38	63387.38	1.167375	0.4735265	0.0204203
Residual	56	3040748.35	54299.08			

**Table S8. Phylogenetic multivariate regression (PGLS) of Von Mises stress MWAM on tooth size (log tooth volume).**

	df	SS.obs	MS	F	P.val	Rsq
log.vol	1	37463.22	37463.22	0.6841096	0.5734266	0.01206881
Residual	56	3066672.51	54762.01			

**Table S9. Phylogenetic multivariate regression (PGLS) of Von Mises stress intervals on tooth size (log tooth Centroid Size).**

	df	SS.obs	MS	F	P.val	Rsq
log.Csize	1	7.235924	7.235924	1.408326	0.5444555	0.02453174
Residual	56	287.725839	5.137961			

**Table S10. Phylogenetic multivariate regression (PGLS) of Von Mises stress intervals on tooth size (log tooth surface area).**

	df	SS.obs	MS	F	P.val	Rsq
log.sa	1	4.866353	4.866353	0.9394005	0.7222777	0.01649825
Residual	56	290.09541	5.180275			

**Table S11. Phylogenetic multivariate regression (PGLS) of Von Mises stress intervals on tooth size (log tooth volume).**

	df	SS.obs	MS	F	P.val	Rsq
log.vol	1	4.012408	4.012408	0.7722816	0.7542458	0.01360315
Residual	56	290.949356	5.195524			



**Table S12. Phylogenetic multivariate regression (PGLS) of Procrustes coordinates on tooth size (log tooth Centroid Size).**

	<b>Df</b>	<b>SS</b>	<b>MS</b>	<b>Rsq</b>	<b>F</b>	<b>Z</b>	<b>Pr(&gt;F)</b>
<b>log.Csize</b>	1	0.002234	0.00223406	0.03928	2.2896	1.7286	0.044*
<b>Residuals</b>	56	0.054641	0.00097573	0.96072			
<b>Total</b>	57	0.056875					

**Table S13. Phylogenetic multivariate regression (PGLS) of Procrustes coordinates on tooth size (log tooth surface area).**

	<b>Df</b>	<b>SS</b>	<b>MS</b>	<b>Rsq</b>	<b>F</b>	<b>Z</b>	<b>Pr(&gt;F)</b>
<b>log.sa</b>	1	0.001997	0.00199688	0.03511	2.0377	1.5029	0.066
<b>Residuals</b>	56	0.054878	0.00097997	0.96489			
<b>Total</b>	57	0.056875					

**Table S14. Phylogenetic multivariate regression (PGLS) of Procrustes coordinates on tooth size (log tooth volume).**

	<b>Df</b>	<b>SS</b>	<b>MS</b>	<b>Rsq</b>	<b>F</b>	<b>Z</b>	<b>Pr(&gt;F)</b>
<b>log.vol</b>	1	0.00186	0.00185955	0.0327	1.8928	1.4	0.084
<b>Residuals</b>	56	0.055015	0.00098242	0.9673			
<b>Total</b>	57	0.056875					

**Table S15. Procrustes ANOVA comparing tooth Procrustes coordinates across dietary categories.**

	<b>Df</b>	<b>SS</b>	<b>MS</b>	<b>Rsq</b>	<b>F</b>	<b>Z</b>	<b>Pr(&lt;F)</b>
<b>diets</b>	4	1.0842	0.27105	0.35224	5.7097	3.6162	0.001**
<b>Residuals</b>	42	1.9938	0.047472	0.64776			
<b>Total</b>	46	3.078					

**Table S16. Phylogenetic Procrustes ANOVA comparing tooth Procrustes coordinates across dietary categories.**

	<b>Df</b>	<b>SS</b>	<b>MS</b>	<b>Rsq</b>	<b>F</b>	<b>Z</b>	<b>Pr(&lt;F)</b>
<b>diets</b>	4	0.005516	0.0013789	0.16908	2.1367	2.0854	0.016*
<b>Residuals</b>	42	0.027105	0.00064535	0.83092			
<b>Total</b>	46	0.03262					

**Table S17. Pairwise PERMANOVA of biomechanical and morphological PC coordinates that explain 90% of the variance, dividing the extant sample in five dietary categories.**

Values indicate Bonferroni corrected p-values. Statistically significant values are marked in bold.

<b>Biomechanical (2 PCs)</b>					
	Omnivore	Herbivore	Insectivore	Durophage	Carnivore
Omnivore		0.366	1	1	<b>0.002</b>
Herbivore	0.366		0.309	1	<b>0.029</b>
Insectivore	1	0.309		0.412	<b>0.001</b>
Durophage	1	1	0.412		0.586
Carnivore	<b>0.002</b>	<b>0.029</b>	<b>0.001</b>	0.586	
<b>Morphological (5 PCs)</b>					
	Omnivore	Herbivore	Insectivore	Durophage	Carnivore
Omnivore		1	1	1	<b>0.001</b>
Herbivore	1		1	1	<b>0.001</b>
Insectivore	1	1		1	<b>0.001</b>
Durophage	1	1	1		0.076
Carnivore	<b>0.001</b>	<b>0.001</b>	<b>0.001</b>	0.076	

**Table S18. Pairwise PERMANOVA of biomechanical and morphological PC coordinates that explain 90% of the variance, dividing the extant sample in three dietary categories.**

Values indicate Bonferroni corrected p-values. Statistically significant values are marked in bold.

<b>Biomechanical (2 PCs)</b>			
	Omnivore	Herbivore	Carnivore
Omnivore		<b>0.0135</b>	<b>0.0003</b>
Herbivore	<b>0.0135</b>		<b>0.0117</b>
Carnivore	<b>0.0003</b>	<b>0.0117</b>	
<b>Morphological (5 PCs)</b>			
	Omnivore	Herbivore	Carnivore
Omnivore		1	<b>0.0003</b>
Herbivore	1		<b>0.0015</b>
Carnivore	<b>0.0003</b>	<b>0.0015</b>	

**Table S19. Accuracy and Kohen’s Kappa comparisons of nine machine learning algorithms tested on the biomechanical data.**

<b>Accuracy</b>						
	<b>Minimum</b>	<b>1st Quartile</b>	<b>Median</b>	<b>Mean</b>	<b>3rd Quartile</b>	<b>Maximum</b>
<b>CART</b>	0.111	0.556	0.667	0.627	0.667	0.889
<b>LDA</b>	0.444	0.667	0.667	0.710	0.778	1.000
<b>MDA</b>	0.222	0.667	0.667	0.687	0.778	1.000
<b>SVML</b>	0.111	0.444	0.667	0.594	0.778	0.889
<b>SVMR</b>	0.111	0.333	0.444	0.463	0.556	0.778
<b>KNN</b>	0.000	0.667	0.667	0.662	0.778	1.000
<b>NB</b>	0.333	0.667	0.778	<b>0.739</b>	0.778	1.000
<b>RF</b>	0.333	0.556	0.667	0.632	0.667	0.889
<b>NNet</b>	0.444	0.667	0.778	<b>0.752</b>	0.778	1.000
<b>Kappa</b>						
	<b>Minimum</b>	<b>1st Quartile</b>	<b>Median</b>	<b>Mean</b>	<b>3rd Quartile</b>	<b>Maximum</b>
<b>CART</b>	-0.400	0.200	0.357	0.326	0.438	0.813
<b>LDA</b>	-0.154	0.357	0.438	0.471	0.600	1.000
<b>MDA</b>	-0.200	0.333	0.400	0.436	0.600	1.000
<b>SVML</b>	-0.500	0.143	0.308	0.324	0.571	0.813
<b>SVMR</b>	-0.500	-0.070	0.118	0.086	0.250	0.625
<b>KNN</b>	-0.500	0.308	0.308	0.362	0.571	1.000
<b>NB</b>	-0.286	0.357	0.571	<b>0.528</b>	0.647	1.000
<b>RF</b>	-0.125	0.250	0.400	0.381	0.507	0.824
<b>NNet</b>	-0.071	0.357	0.571	<b>0.524</b>	0.600	1.000

**Table S20. Accuracy and Kohen’s Kappa comparisons of nine machine learning algorithms tested on the morphological data.**

<b>Accuracy</b>						
	<b>Min.</b>	<b>1st Q</b>	<b>Median</b>	<b>Mean</b>	<b>3rd Q</b>	<b>Max.</b>
<b>CART</b>	0.222	0.556	0.667	0.636	0.778	0.889
<b>LDA</b>	0.111	0.444	0.556	0.577	0.667	0.889
<b>MDA</b>	0.333	0.556	0.667	0.668	0.778	1.000
<b>SVML</b>	0.222	0.444	0.556	0.564	0.667	0.889
<b>SVMR</b>	0.222	0.667	0.667	0.693	0.778	1.000
<b>KNN</b>	0.222	0.556	0.667	0.674	0.778	0.889
<b>NB</b>	0.222	0.444	0.556	0.586	0.667	0.889
<b>RF</b>	0.333	0.667	0.667	0.703	0.778	1.000
<b>NNet</b>	0.333	0.667	0.667	<b>0.708</b>	0.778	1.000
<b>Kappa</b>						
	<b>Min.</b>	<b>1st Q</b>	<b>Median</b>	<b>Mean</b>	<b>3rd Q</b>	<b>Max.</b>
<b>CART</b>	-0.286	0.200	0.345	0.351	0.571	0.800
<b>LDA</b>	-0.333	0.143	0.333	0.298	0.471	0.824
<b>MDA</b>	-0.125	0.283	0.471	0.442	0.625	1.000
<b>SVML</b>	-0.400	0.118	0.250	0.283	0.400	0.824
<b>SVMR</b>	-0.313	0.308	0.419	0.435	0.600	1.000
<b>KNN</b>	-0.125	0.283	0.400	0.418	0.571	0.824
<b>NB</b>	-0.400	0.118	0.294	0.287	0.446	0.824
<b>RF</b>	-0.125	0.308	0.400	0.461	0.600	1.000
<b>NNet</b>	0.000	0.383	0.498	<b>0.501</b>	0.625	1.000

**Table S21. Diet prediction and dietary class probability of early dinosaurs.** The classifications are based on the final machine learning model for the biomechanical and morphological data.

Species	Prediction	Carnivore probability	Herbivore probability	Omnivore probability
<b>Biomechanical classification</b>				
<i>Manidens condorensis</i>	Carnivore	0.953	0.046	0.001
<i>Lesothosaurus diagnosticus</i>	Omnivore	0.001	0.093	0.906
<i>Herrerasaurus ischigualastensis</i>	Carnivore	0.861	0.138	0.001
<i>Eodromaeus murphi</i>	Carnivore	0.997	0.002	0.001
<i>Buriolestes schultzi</i>	Carnivore	0.574	0.425	0.001
<i>Eoraptor lunensis</i>	Carnivore	0.959	0.040	0.001
<i>Saturnalia tupiniquim</i>	Omnivore	0.006	0.350	0.644
<i>Thecodontosaurus antiquus</i>	Herbivore	0.218	0.782	0.000
<i>Plateosaurus engelhardti</i>	Herbivore	0.010	0.545	0.445
<i>Massospondylus carinatus</i>	Herbivore	0.022	0.945	0.034
<i>Ngwevu intloko</i>	Herbivore	0.024	0.972	0.004
<b>Morphological classification</b>				
<i>Manidens condorensis</i>	Herbivore	0.004	0.996	0.000
<i>Lesothosaurus diagnosticus</i>	Herbivore	0.016	0.919	0.065
<i>Herrerasaurus ischigualastensis</i>	Carnivore	0.954	0.040	0.006
<i>Eodromaeus murphi</i>	Carnivore	0.958	0.018	0.024
<i>Buriolestes schultzi</i>	Carnivore	0.971	0.001	0.028
<i>Eoraptor lunensis</i>	Carnivore	0.586	0.412	0.002
<i>Saturnalia tupiniquim</i>	Carnivore	0.855	0.123	0.022
<i>Thecodontosaurus antiquus</i>	Herbivore	0.126	0.871	0.003
<i>Plateosaurus engelhardti</i>	Herbivore	0.013	0.987	0.000
<i>Massospondylus carinatus</i>	Herbivore	0.036	0.963	0.001
<i>Ngwevu intloko</i>	Herbivore	0.020	0.980	0.000

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