



## An exceptionally well-preserved skeleton of *Thomashuxleya externa* (Mammalia, Notoungulata), from the Eocene of Patagonia, Argentina

Juan D. Carrillo and Robert J. Asher

### ABSTRACT

We describe one of the oldest notoungulate skeletons with associated craniodental and postcranial elements: *Thomashuxleya externa* (Isotemnidae) from Cañadón Vaca in Patagonia, Argentina (Vacan subage of the Casamayoran SALMA, middle Eocene). We provide body mass estimates given by different elements of the skeleton, describe the bone histology, and study its phylogenetic position. We note differences in the scapulae, humerii, ulnae, and radii of the new specimen in comparison with other specimens previously referred to this taxon. We estimate a body mass of  $84 \pm 24.2$  kg, showing that notoungulates had acquired a large body mass by the middle Eocene. Bone histology shows that the new specimen was skeletally mature. The new material supports the placement of *Thomashuxleya* as an early, divergent member of Toxodontia. Among placentals, our phylogenetic analysis of a combined DNA, collagen, and morphology matrix favor only a limited number of possible phylogenetic relationships, but cannot yet arbitrate between potential affinities with Afrotheria or Laurasiatheria. With no constraint, maximum parsimony supports *Thomashuxleya* and *Carodnia* with Afrotheria. With Notoungulata and Litopterna constrained as monophyletic (including *Macrauchenia* and *Toxodon* known for collagens), these clades are reconstructed on the stem to Euungulata (i.e., Perissodactyla and Artiodactyla). Unconstrained, Bayesian analysis weakly supports the possibility that *Thomashuxleya* is a stem xenarthran; with Notoungulata and Litopterna constrained as monophyletic, the two clades are recovered as sister to Perissodactyla. Anatomical data sampled thus far for *Thomashuxleya*, combined with collagen amino acids for Pleistocene meridiungulates, substantially limit the number of possible affinities for endemic South American species among mammals, although ambiguity still remains.

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## INTRODUCTION

South America was isolated during most of the Cenozoic and was home to a highly endemic fauna (Simpson, 1980; Pascual, 2006; Wilf et al., 2013). South American Native Ungulates, or “SANUs” (Welker et al., 2015), were classified in Meridiungulata by McKenna (1975). Meridiungulates are a conspicuous faunal element of the South American Cenozoic, with an extensive fossil record that spans the early Paleocene (~ 64 Ma, Tiupampan South American Land Mammal age or SALMA; Gelfo et al., 2009; Woodburne et al., 2014a, 2014b) to late Pleistocene (~11-7 ka, Cione et al., 2003; ~ 11-13 ka, Barnosky and Lindsey, 2010). Notoungulata is the major clade within Meridiungulata and exhibits a high taxonomic diversity (> 140 genera and 13 families; Croft, 1999), large morphological disparity, wide range of body masses (Giannini and García-López, 2014), different degrees of hypsodonty, and diverse diets (MacFadden, 2005; Townsend and Croft, 2008; Cassini et al., 2011; Madden, 2015). The monophyly of Notoungulata is generally accepted, with Pyrotheria sometimes included (Patterson, 1977; Billet, 2010, 2011) or excluded (Roth, 1903; Cifelli, 1993; Simpson, 1978). Alignments of amino acid residues of alpha 1 and 2 collagen chains support a close relationship of Pleistocene members of Notoungulata (*Toxodon*) and Litopterna (*Macrauchenia*) with extant Perissodactyla (Welker et al., 2015). Another analysis of alpha 1 and 2 collagen amino acids also supported *Toxodon* and *Macrauchenia* with perissodactyls. However, this study (Buckley, 2015, figure 2) gave likelihood bootstrap support values below 50 for this clade, and (unlike Welker et al., 2015) the alignments from Buckley (2015) are not publicly accessible as of this writing.

Most of the earliest members of the notoungulate radiation in South America are known from dental and isolated postcranial remains (e.g., Horowitz, 2004; Bergqvist, et al., 2007; Shockley, and Flynn, 2007; Lorente et al., 2014; Lorente, 2015). Specimens with associated cranial and postcranial elements are extremely rare. The Isotemnidae (Notoungulata) has been interpreted as a basal group within Toxodontia (Simpson, 1936; Cifelli, 1993). More recent evidence suggests that Isotemnidae could be polyphyletic (Billet, 2011) and defined by plesiomorphic traits among Toxodontia (Simpson, 1967; Billet, 2011).

Remains of *Thomashuxleya* (Isotemnidae) are among the earliest known associated notoungulate skeletons (Simpson, 1936, 1967). However,

the original descriptions of Simpson were idealized and based on a composite skeleton representing three different genera: *Thomashuxleya*, *Anisotemnus*, and *Pleurostyloodon* (Simpson, 1967; Shockley and Flynn, 2007). This reduces the value of Simpson's skeletal reconstruction for palaeobiological inference in general and phylogenetic analysis in particular. *Thomashuxleya* is middle Eocene in age, and our material derives from the Vacan subage of the Casamayoran SALMA, older than the adjacent Barrancan subage (Cifelli, 1985; Gelfo et al., 2009; Kay et al., 1999; Woodburne et al., 2014a). *Thomashuxleya* has previously been referred to a relatively basal clade within Toxodontia (Billet, 2011), making it particularly relevant to understand the early radiation of notoungulate mammals.

At least 28 mammal taxa are recognized for the Cañadón Vaca local fauna (Cifelli, 1985, table 5; Shockley and Flynn, 2007). Among Isotemnidae (*sensu* McKenna and Bell, 1997), four species are present: *Pleurostyloodon similis*, *Isotemnus primitivus*, *Thomashuxleya externa*, and *Anisotemnus distortus* (Cifelli, 1985; Shockley and Flynn, 2007).

In this contribution, we describe a specimen consisting of a single individual of *Thomashuxleya externa* with a well-preserved skull and jaws associated with postcrania from Cañadón Vaca, east of Colhué Huapí Lake in Chubut Province, Argentina (Figure 1) (Simpson, 1948; Cifelli, 1985). Parts of most elements of the skeleton are represented, including the skull, mandible, vertebrae, fore- and hind-limbs, shoulder, and pelvic girdles. Associated remains of other individuals found at the same locality include a partially articulated manus. This discovery provides an unusually complete anatomical basis to study the biology and phylogenetic position of an early member of toxodont notoungulates.

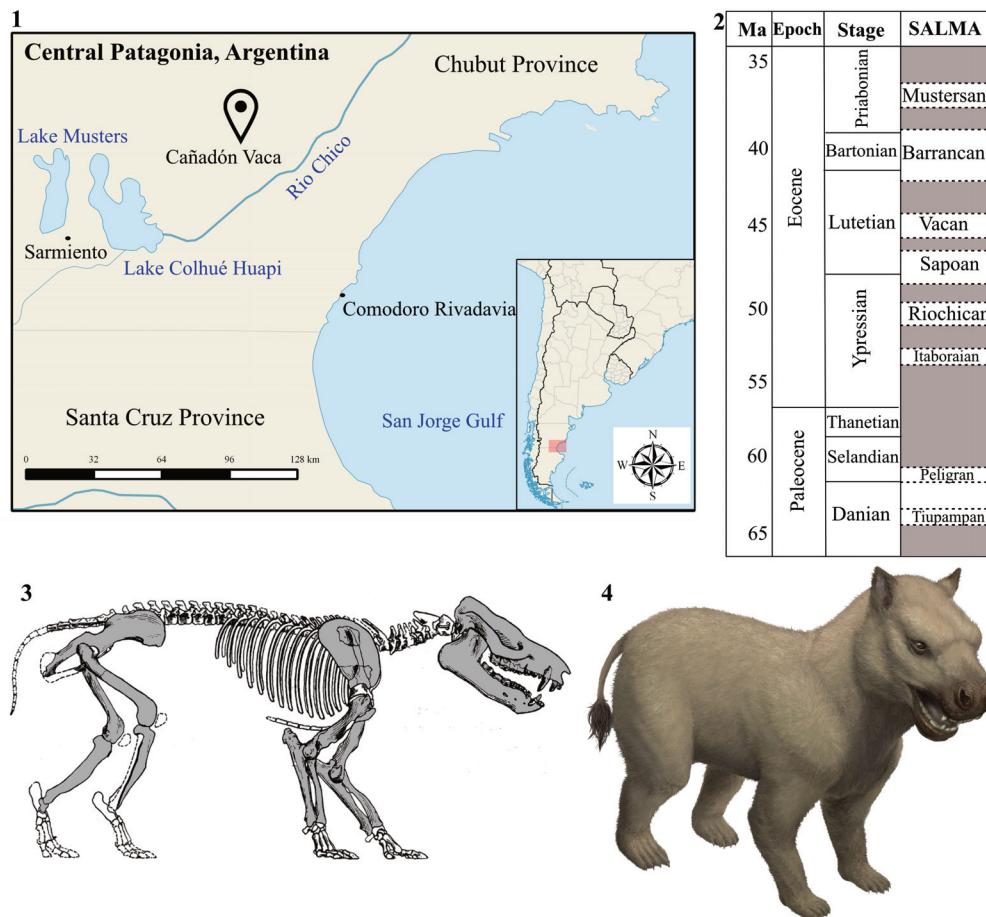
## MATERIALS AND METHODS

### Institutional Abbreviations

MPEF-PV: Museo Paleontológico Egidio Feruglio-Paleontología de Vertebrados, Trelew, Chubut, Argentina; AMNH: American Museum of Natural History, New York, USA; MNHN: Muséum national d'Histoire naturelle, Paris, France.

### Anatomical Description

We follow Smith and Dodson (2003) for the dental orientation, where mesial and distal designate the tooth surface directions facing forward and away from the mandibular symphysis, respec-



**FIGURE 1.** Geographical and stratigraphical occurrence of MPEF-PV 8166. **1** location of Cañadón Vaca, Chubut, Argentina; **2** Paleogene time table and South American Land Mammal Ages (SALMAs) after (Woodburne et al. 2014a,b); **3** skeletal restoration of *Thomashuxleya* modified from Simpson (1936); **4** artistic reconstruction of *Thomashuxleya externa* (by Stjepan Lukac).

tively. We follow Madden (1990) for the notoungulate dental terminology and Cerdeño et al. (2012) for the orientation of the postcranial bones.

The skull, mandible, and dentition of MPEF-PV 8166 were compared with different isotemnids as reviewed by Simpson (1967) and the original descriptions of Ameghino (1901). The postcranial elements of MPEF-PV 8166 were compared with AMNH 28905 referred to *T. externa* and originally described by Simpson (1936, 1967), and recently reviewed by Shockley and Flynn (2007). In addition, O'Leary et al. (2013) included much of the anatomy of published specimens in their phylogenetic study of living and fossil mammals. We made additional comparisons with other isotemnid specimens as described by Simpson (1936, 1967) and reviewed by Shockley and Flynn (2007), and also with other basal notoungulates (Bergqvist et al., 2007; Lorente et al., 2014; Lorente, 2015). Our associated remains of *Thomashuxleya* (primarily

MPEF-PV 8166) provided the basis for revising and scoring new postcranial character states for *T. externa* using the morphological dataset of O'Leary et al. (2013). Our revised character states have been accessioned in MorphoBank (Project 2084).

#### Body Mass Estimation

We used the specimen to test the congruence between body mass estimations with regression equations from the crania and postcrania. We took standard linear measurements of the limb bones, skull, and dentition with calipers to the nearest 0.1 mm. When the right and left limb bones could be measured, we used the average. In order to generate body mass estimations for MPEF-PV 8166, we used craniodental regression equations selected from Janis (1990) and multivariate regression algorithms following Mendoza et al. (2006). Postcranial regression equations follow Scott (1990) for the

limb bones and Tsubamoto (2014) for the astragalus (see Appendix 1-2).

The regression equations were selected based on their high  $r^2$  value ( $> 0.90$ ), a percentage of predicted error (PE) below 40%, and presence of relevant anatomy in MPEF-PV 8166. Janis (1990) and Scott (1990) obtained regression equations using a dataset of extant ungulates and these have been used previously to estimate body mass for notoungulates (e.g., Cassini et al., 2012; Elis-samburu, 2012). The dataset of Tsubamoto (2014) included several orders of living mammals.

To obtain body mass estimates we log-transformed (base 10) the body mass estimates from the regression equations from Janis (1990) and Scott (1990), and used a natural log transformation for Tsubamoto (2014). We compared the results from the different subsets of measurements, namely skull, teeth, long bones, and astragalus. We calculated the statistical descriptors and moments (minimum, maximum and mean values, median, mode, skewness, and kurtosis). We resampled with replacement (i.e., bootstrap) with 1000 replications and calculated 95% confidence intervals (CI) using R (R Core Team, 2016).

### Bone Histology

We took a cortical slice of 1 cm from the mid-shaft of the right femur. The section was embedded in Araldite® 2020 prior to sawing and grinding. For the production of thin sections we followed protocols of Straehl et al. (2013). The section (PHZ 950) was observed in normal transmitted and cross-polarized light using a Leica DM 2500 M microscope equipped with a Leica DFC 420 C digital camera. In order to infer the bone microstructure, we also took a photograph of the complete section and transformed to a binary image with Affinity Designer® 1.5.4, where black represents the bone and white the cavities.

### Phylogenetic Analysis

**Relationships of *Thomashuxleya* within Notoungulata.** To study the phylogenetic relationship of *Thomashuxleya* within Isotemnidae, and with other Paleogene notoungulates, we added the information of the new *Thomashuxleya* specimen (MPEF-PV 8166) to the character matrix of Deraco and García-López (2015). The matrix includes 68 taxa (59 of which are notoungulates) and 146 craniodental characters. We ordered 10 characters, following Deraco and García-López (2015); two others (37 and 119) ordered by them are binary. Searches used 500 replicates of random

addition sequence, holding 10 trees per replication, using tree bisection reconnection (TBR) for branch swapping.

**Relationships of *Thomashuxleya* within Placentalia.** Postcranial characters were scored for the almost complete *Thomashuxleya externa* specimen (MPEF-PV 8166), using the morphological dataset of O'Leary et al. (2013; Morphobank project 773); cranial characters were the same as in O'Leary et al. (2013). The specimens used by O'Leary et al. (2013) to score the craniodental characters were referred to *T. externa* by Simpson (1967) and are not associated with postcranial remains. This resulted in 3660 craniodental and postcranial characters for 87 taxa, available in MorphoBank (Project 2084). We concatenated the morphological dataset with the amino acid alignment of Meredith et al. (2011; TreeBase number S11872), which comprises 11010 amino acids for 169 taxa, and the collagen alignment of Welker et al. (2015), with 2028 amino acids for 77 taxa. For the phylogenetic analysis, we combined the morphological and amino acid data using R (R Core Team, 2016), merging species of the same genus and where necessary coding data as absent for fossils (e.g., for most sequence data). To increase the analytical tractability of our dataset, we excluded non-mammals, living mammals with more than 50% missing data, non-placental fossils, and fossils with over 90% missing data (except meridiungulates) from our morphological dataset.

For the parsimony analysis, we applied parsimony as the optimality criterion in TNT (Goloboff et al., 2008a), PAUP 4.0a150 (Swofford, 2002), and PAUPr (Sykes and Lewis, 2001) to a dataset consisting of 182 taxa and 16698 characters (13038 amino acids and 3660 morphological characters). All characters were unordered. We explored equal weights and extended implied weighting; the latter weights the two character sets (amino acids and morphology) using their average homoplasy (Goloboff et al., 2008b, Goloboff, 2014). In order to account for characters with many missing entries that would receive artificially high implied weights, we used the command "xpiwe (\*", which sets a concavity value (k) according to the number of missing entries (Goloboff, 2014). In addition to our unconstrained analyses, we also explored a number of backbone constraints. Following Tarver et al. (2016) we constrained monophyly of Atlantogenata (Xenarthra + Afrotheria) and Boreutheria (Laurasiatheria + Euarchontoglires) as shown in Tarver et al. (2016, figure 1). We also explored the impact of constraining 1) monophyly

of each meridiungulate clade in our sample (i.e., the two notoungulates and two litopterns) and 2) all six meridiungulates. We compared the resulting trees of the unconstrained and constrained analyses using a Wilcoxon rank sum test in PAUP\* (Swofford, 2002). Parsimony searches included 500 replicates of random addition sequence, holding 10 trees per replication, using tree bisection reconnection (TBR) for branch swapping.

To make Bayesian searches tractable given our available computing resources, we further decreased the number of living taxa to those under 25% missing data (except meridiungulates), leaving 51 taxa. We obtained partitioning schemes and substitution models for the protein alignment with Partition Finder Protein v1.1.0 using the strict hierarchical clustering algorithm and the Bayesian Information Criterion for model selection (Lanfear et al., 2012). The best model partition scheme resulted in 20 partitions (out of 22 total coding genes) and assigned the JTT+G model for each amino acid partition and the MTMAM+G model for the collagen alpha 1 and 2. We used the standard discrete model implemented by MrBayes (Ronquist et al. 2011) for the morphological characters (Table 1). The dataset was analysed with MrBayes (Ron-

quist et al., 2012) and BEAGLE to utilize both GPU and CPU during searches. We used Tracer (Rambaut et al., 2014) for the visualisation and diagnostics of the MCMC output. We ran two analyses, one unconstrained and one constraining monophyly of the two notoungulates and two litopterns in our sample, as described above.

We used three runs of 3,000,000 generations with five chains (four heated and one cold) for the unconstrained analysis and two runs of 3,500,000 generations with four chains (three heated and one cold) for the constrained analysis. We sampled every 1000 generations, and use a temperature of 0.5.

## RESULTS

### Systematic Paleontology

NOTOUNGULATA Roth, 1903

ISOTEMNIDAE Ameghino, 1897

*Thomashuxleya* Ameghino, 1901

**Type species.** *Thomashuxleya rostrata* Ameghino, 1901, by original designation

*Thomashuxleya externa* (Ameghino, 1901)

*Thomashuxleya artuata* Ameghino, 1901;

**TABLE 1.** Best partition scheme for Bayesian analysis as obtained from Partition Finder (Lanfear et al., 2012).

Best model	Subset partitions	Subset sites
JTT+G	TTN	1-1479
JTT+G	CNR1	1480-1814
JTT+G	BCHE	1815-2146
JTT+G	EDG1	2147-2466
JTT+G	RAG1	2467-3066
JTT+G	RAG2	3067-3215
JTT+G	ATP7A	3216-3444
JTT+G	TYR1	3445-3587
JTT+G	Adora3	3588-3698
JTT+G	BDNF	3699-3885
JTT+G	ADRB2	3886-4153
JTT+G	PNO	4154-4260
JTT+G	A2AB	4261-4543
JTT+G	BRCA1, BRCA2	4544-5574, 5575-7256
JTT+G	APOB, DMP1	7257-7710, 9718-10592
JTT+G	GHR	7711-8026
JTT+G	VWF	8027-8417
JTT+G	ENAM	8418-9717
JTT+G	IRBP	10593-11010
MTMAM+G	Collagen alpha 1 and 2	11011-13038
Standard discrete	Morphology	13039-16698

**TABLE 2.** Skeletal elements of isotemnid specimens from Cañadón Vaca.

	<i>T. externa</i> MPEF-PV 8166	<i>T. externa</i> MNHN CAS 844	<i>T. externa</i> MNHN CAS 64	<i>T. externa</i> AMNH 28698	<i>T. externa</i> AMNH 28447	<i>T. externa</i> AMNH 28905	<i>T. externa</i> AMNH 28563	<i>cf. T. externa</i> AMNH 142463	<i>A. distentus</i> AMNH 28906	<i>A. distentus</i> AMNH 28647	<i>P. similis</i> AMNH 28904	<i>P. similis</i> AMNH 28635	Isotemnidae indet. AMNH 28690
Skull	x	x		x							M1 only		
Mandible	x		x		x								
Scapula	x				x			x					
Humerus	x				x	x		x		x			
Ulna	x				x	x		x		x			
Radius	x					x		x	x	x			
Manus								x	x	x			
Pelvis	x				x							x	x
Femur	x				x						x	x	x
Tibia	x								x	x	x	x	x
Pes	x					x	x		x		x	x	

synonymized with *T. externa* by Simpson, 1967, p. 159

**Stratigraphic provenance.** MPEF-PV 8166 came from the Cañadón Vaca member of the Sarmiento Formation (Bellosi and Krause, 2014). An age of ~45 Ma has been proposed for the “Vacan” subage (Cifelli, 1985; Carlini et al., 2005; Woodburne et al., 2014). Bellosi and Krause (2014) estimated a range of 43.1–46.9 Ma for the Cañadón Vaca member.

**Thomashuxleya systematics.** Ameghino (1901) first described *Thomashuxleya* based on material from the “couches a *Notostylops*” (now known to be part of the Casamayoran, middle Eocene) and originally assigned it to Homalodotheriidae (“Homalodontotheridae” in Ameghino, 1901). Ameghino designated *T. rostrata* as the type species (Ameghino, 1901, p. 177). In the same publication, Ameghino recognized three more species within the genus: *T. artuata*, *T. robusta*, and *T. externa* (Ameghino, 1901, p. 179).

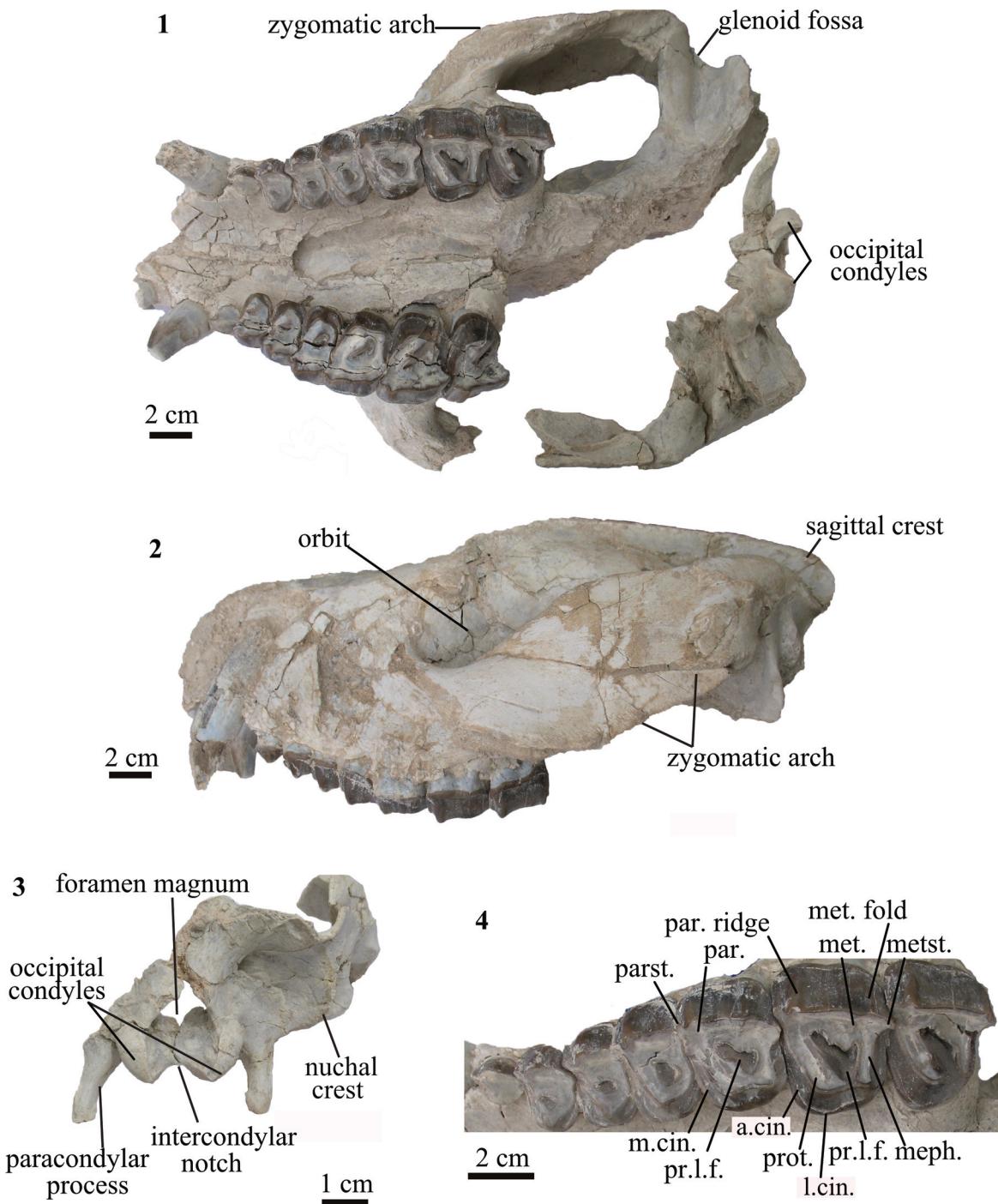
Simpson (1967) considered *Thomashuxleya* to be part of Isotemnidae. Simpson (1967, p. 121) considered five Casamayoran Isotemnidae genera to be valid: *Pleurostyloodon* (Ameghino, 1897), *Anisotemnus* (Ameghino, 1902), *Plexotemnus* (Ameghino, 1904a), *Isotemnus* (Ameghino, 1897), and *Thomashuxleya* (Ameghino, 1897). For *Thomashuxleya*, Simpson (1967) recognized two species: *T. externa* and *T. rostrata*; the latter is recorded in the local fauna south of Colhué Huapí Lake in Gran Barranca, (Barracan; Woodburne et al., 2014a). Other isotemnids present in the fauna

of Colhué Huapí are *Pleurostyloodon modicus* and *Isotemnus primitivus* (Cifelli, 1985).

Shockley and Flynn (2007) studied several of the postcranial elements referred to *Thomashuxleya* by Simpson (1936, 1967), and recognized that at least three genera were included in Simpson’s (1936, 1967) skeletal reconstruction. Based on the material then available, they inferred a straight posture, plantigrade or semi-digitigrade locomotion, and lack of running among Vacan isotemnids. Of the known Isotemnidae specimens recovered from Cañadón Vaca, MPEF-PV 8166 is the only one with associated craniodental and postcranial elements (Table 2).

### Description

**Skull.** MPEF-PV 8166 preserves most of the skull, including the cranium with maxillae, palatines, zygomatic arches, nasal, frontal, temporal and occipital, and the almost complete mandible. The skull of MPEF-PV 8166 measures 28 cm from the most posterior point of the sagittal crest to the most anterior point of the snout. Due to the preservation, the sutures are not visible. In lateral view, the rostrum is high (Figure 2.2). The zygomatic arch is wide and robust, more than in *Pleurostyloodon* (Simpson, 1967). There is a well-defined sagittal crest (Figure 2.2), and the glenoid fossa is wide and concave (Figure 2.1), as in *Pleurostyloodon* (Simpson, 1967). The occipital condyles are oval with a deep intercondylar notch. The paracondylar processes of the exoccipitals are long and narrow. The nuchal crest forms a semicircular outline in



**FIGURE 2.** Skull of *T. externa* (MPEF-PV 8166). **1** ventral view; **2** lateral view; **3** occiput in caudal view; **4** detail of upper dentition in occlusal view. Abbreviations are prot=protocone (part of the protoloph), par=paracone (part of the ectoloph), parst=parastyle, met=metacone (part of the ectoloph), metst=metastyle, meph=metaloph, pr.l.f.=primary lingual fold, m.cin=mesial cingulum, l.cin=lingual cingulum.

posterior view and does not project below the level of the occipital condyles (Figure 2.3). The palate is triangular (Figure 2.2).

**Upper dentition.** The upper canines are clearly differentiated from the rest of the teeth. The mesiodistal length of each tooth steadily increases from P1 to M3 (Figure 2.1). The P1 is longer than wide, in contrast with *Anisotemnus distentus*, which has the opposite condition (Simpson, 1967). The P1 is oval in occlusal view and with a single mediolabial cusp (Figure 2.4). It has no mesial or distal stylar projections, in contrast to *Pleurostyłodon* (Simpson, 1967). The P2-P4 are similar, increasing in labio-lingual width from P2 to P4 (Figure 2.4). They have a well-defined paracone ridge, and mesial to it there is a small parastyle, less prominent than in *Pleurolystodon* (Simpson, 1967). The metacone fold is absent (Figure 2.4). There is a mesial and distal cingulum, as in *Pleurolystodon* (Simpson, 1967).

The upper molars have a paracone ridge (Figure 2.4) which is less developed than in *Pleurostyłodon* (Simpson, 1967). The metacone fold is observed as a very smooth ridge (Figure 2.4), less defined than in *Pleurostyłodon* and *Isotemnus* (Simpson, 1967). There is a metastyle, which is more defined in M2 and M3 than in M1 (Figure 2.4). The upper molars have a primary lingual fold surrounded by the protoloph, ectoloph, and metaloph (Figure 2.4). As the teeth wear down, the primary lingual fold closes, forming a fossa (Simpson, 1967), as seen in M1 (Figure 2.4). There is a continuous labial, mesial, and lingual cingula, as in *Thomashuxleya* (Simpson, 1967). The mesial cingulum surrounds the protocone, reaching the lingual face of the teeth (Figure 2.4), in contrast to *Isotemnus*, which does not have lingual cingula (Simpson, 1967). The upper molars have no fossettes, which were "numerous and somewhat persistent with wear" in *Anisotemnus* according to Simpson (1967, p. 136). MPEF-PV 8166 is referred to *Thomashuxleya* based on the presence of a continuous lingual cingulum in the upper molars and dental dimensions (Simpson, 1967).

In addition of the differences with *Pleurostyłodon* and *Isotemnus* mentioned above, MPEF-PV 8166 is about 60% larger than the specimens referred to *P. similis* and about 85% larger than *I. primitivus* (also recorded in Cañadón Vaca) based on the upper molar dimensions (Table 3; Simpson, 1967, table 45,49,54).

**Mandible.** The horizontal ramus is straight (Figure 3.2). The vertical ramus is wide and high (Figure 3.2). The masseteric fossa is large and oval (Fig-

ure 3.2). The condylar process is wide medio-laterally and narrow rostro-caudally (Figure 3.1). The coronoid process is narrow and high, extending more dorsally than the condylar process (Figure 3.2). The symphysis extends caudally to the level of p3/p4 (Figure 3.1), whereas in *Pleurostyłodon* it extends to the level of p2 (Simpson, 1967). There are two mental foramina in the right horizontal ramus, one at the level of the mesial border of p2 and the other at the level of the mesial border of p3. In *Pleurostyłodon*, the mental foramina are at the level of p3 and p4 (Simpson, 1967).

**Lower dentition.** The lower dentition has a small diastema of approximately 2 mm between p1 and p2. As in the upper teeth, the size increases from p1 to m3 (Figure 3.1). The p1 is triangular in lingual and labial views (Figure 3.2), and it has a median main cusp (Figure 3.3). The p2 has a large protoconid, with a crest projecting mesially, as in *Pleurostyłodon* (Simpson, 1967). The metaconid projects lingually (Figure 3.3) and at its base shows a cingulid (Figure 3.2). The p3 and p4 are bicrescentic in occlusal view and show a high degree of wear. They have a well-defined labial fold (Figure 3.3) and a labial cingulid (Figure 3.2).

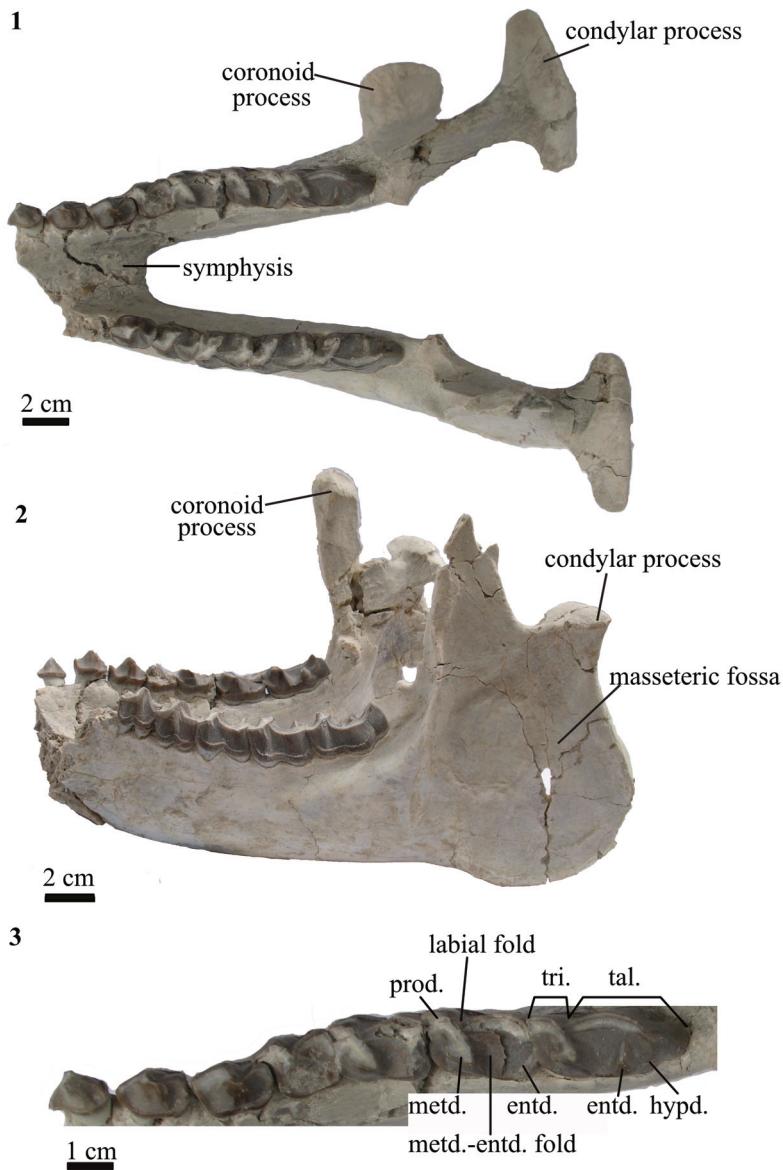
The lower molars have well-defined labial and lingual cingulids (Figure 3.2). The talonids are longer mesio-distally than the trigonids (Figure 3.3). The metaconid projects lingually and not distally as in *Pleurostyłodon* (Simpson, 1967). The meta-entocenid fold is mesio-distally long and labio-lingually broad (Figure 3.3). The m3 has a long talonid, with the entoconid separated from the hypoconulid (Figure 3.3), as in *Pleurostyłodon* (Simpson, 1967). Of the two species currently recognized for *Thomashuxleya*, we assign MPEF-PV 8166 to *T. externa* based on the well-defined labial and lingual cingulid (Ameghino, 1901) and its dental dimensions (Table 3; Simpson, 1967).

**Scapula.** The scapulae of MPEF-PV 8166 differ from AMNH 28905, previously referred to *T. externa* and *Anisotemnus distentus* (AMNH 28906). The comparison between the two *T. externa* specimens is difficult because MPEF-PV 8166 does not preserve the complete blade, and AMNH 28905 has been modified by post-recovery restoration. When comparing MPEF-PV 8166 with AMNH 28905 as coded by (O'Leary et al., 2013), we observed some differences: the coracoid process is pointing towards the axillary edge of the scapula and not perpendicular to the blade (Figure 4.1-2); the infraspinous fossa does not reach the rim of the glenoid fossa; and it is triangular rather than rectangular in lateral view (Figure 4.3).

**TABLE 3.** Dental measurements of *Thomashuxleya*; \* =taken from Simpson (1967).

		p1		p2		p3		p4		m1		m2		m3	
		Length	Width												
<i>T. rostrata</i>	MACN 10370*	13.7	11.8	17.5	12.7	18.1	14.8	20.9	16.6						
	MACN 10539*							19.9	17.8	26	19.8	26.4	19.5	37.2	16.9
	MACN 10546*							21.5	16.8						
	AMNH 28692*					19.5	16.5			26	16.4	28.5	18.8		
	AMNH 28764*							21.8	13.7						
	Mean	13.7	11.8	17.5	12.7	18.8	15.7	21.0	16.2	26	18.1	27.5	19.2	37.2	16.9
	SD	NA	NA	NA	NA	1.0	1.2	0.8	1.8	0.0	2.4	1.5	0.5	NA	NA
	n	1	1	1	1	2	2	4	4	2	2	2	2	1	1
<i>T. externa</i>	MACN 10540*					16.5	14.6	19	17.2	21		28		37	
	MACN 10537*							16.3	13	19.5	14	23	15.5		
	AMNH 28447*			14.2	11.1	16.2	11.6	18.1	13	22.0	15.6	27.5	16.5	31.6	15.8
	AMNH 28697*							17.3	14.2	23.2	15.9	25.5	17.2	36	16.6
	AMNH 28756*			14.7	10.9	15.7	12.8	16.4	14.2	21.8	16.3	25.6	17.8	32.3	16.3
	AMNH 28698*	11.0	10.0	15.1	12.6	16.7	13.6	16.5	14.6	24	16.1	26	17.8	37	17
	AMNH 28686*												34.5	17.0	
	AMNH 28822*									20	16.4	23	18.0		
	Mean*	11	10	14.7	11.5	16.3	13.2	17.3	14.4	21.6	15.7	25.5	17.1	34.7	16.5
	SD*	NA	NA	0.5	0.9	0.4	1.3	1.1	1.5	1.6	0.9	2.0	1.0	2.4	0.5
	n*	1	1	3	3	4	4	6	6	7	6	7	7	6	5
	MPEF-PV 8166	12.8	8.3	15.4	10	16.5	10.1	16.4	13.3	20.1	15.6	22.5	15.3	32.3	15.7

		P1		P2		P3		P4		M1		M2		M3	
		Length	Width												
<i>T. rostrata</i>	MACN 10370*	15.4	15.7	17.0	27.2	19.5	31.2	21	35						
	MACN 10542*									29.5	41.8	33.0	44.2	28.8	40.8
	Mean*	15.4	15.7	17	27.2	19.5	31.2	21	35	29.5	41.8	33	44.2	28.8	40.8
	SD*	NA	NA												
	n*	1	1	1	1	1	1	1	1	1	1	1	1	1	1
<i>T. externa</i>	MACN 10543*									24	34	28.5	35.9		
	AMNH 28699*			17.5	22.1	16.8	26.4	18.5	32.1	23.8	35.8	26.7	42		39.5
	AMNH 28757*	13.8	11.5	15.2	22.1	18	26			25.7		27.2	39.2		
	AMNH 28698*	10	11			16		18	33.2	23.9	38.7	27.2	41	26	41
	MHN CAS 844	10.84	13.86	17.79	22.93	18.94	26.80	19.13	31.25	21.40	32.75	22.71	35.98	26.55	35.46
	Mean*	11.5	12.1	16.8	22.4	17.4	26.4	18.5	32.2	23.8	35.3	26.5	38.8	26.3	38.7
	SD*	2.0	1.5	1.4	0.5	1.3	0.4	0.6	1.0	1.5	2.6	2.2	2.8	0.4	2.9
	n*	3	3	3	3	4	3	3	3	5	4	5	5	2	3
	MPEF-PV 8166	10.2	10.5	14.0	16.2	15.0	21.6	14.6	25.9	25.2	31.3	27.9	36.9	24.0	32.0

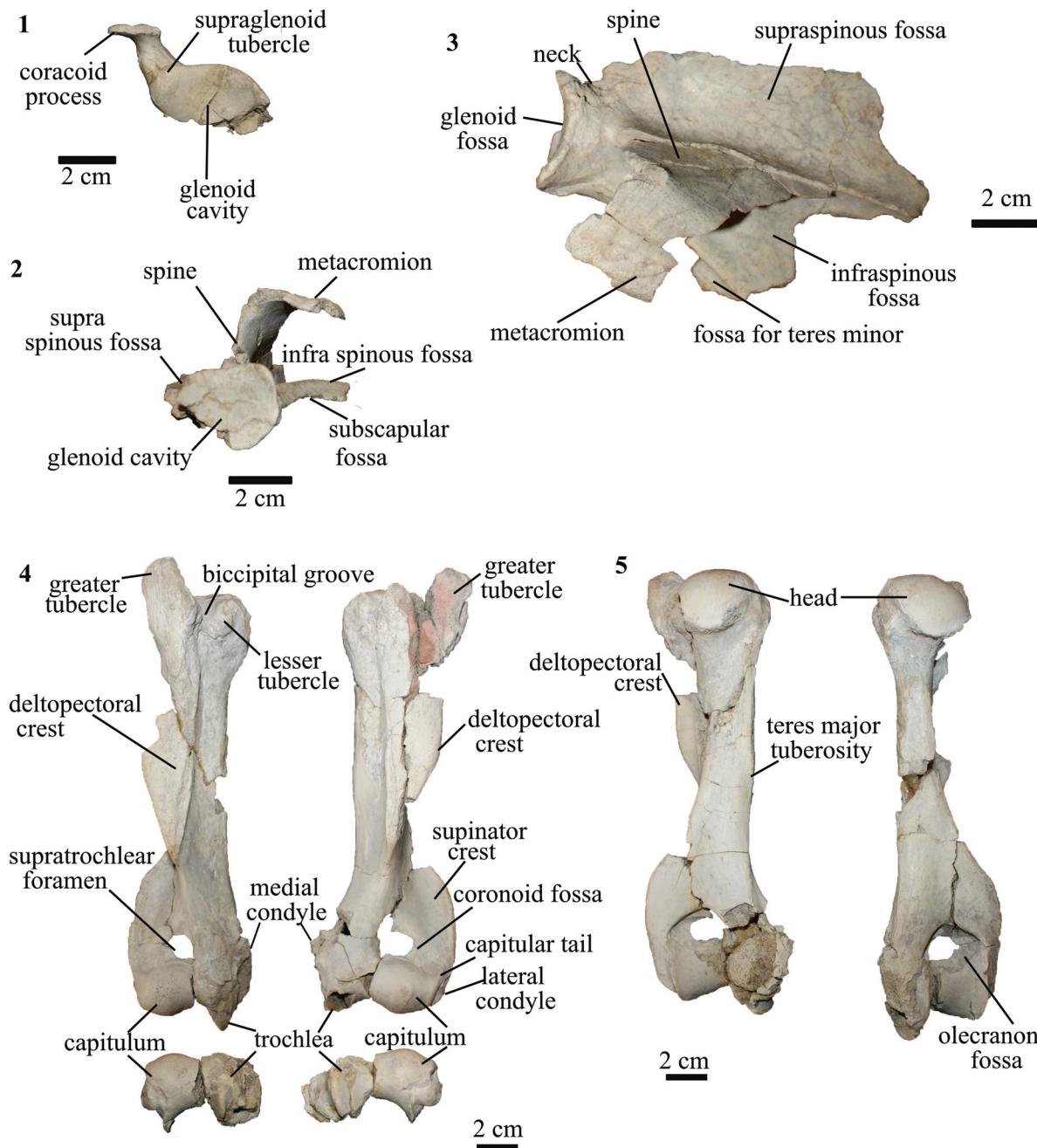


**FIGURE 3.** Mandible of *T. externa* (MPEF-PV 8166) in **1** dorsal and **2** lateral views; **3** lower dentition in occlusal view. Abbreviations are tri=trigonid, tal=talonid, prod=protoconid, metd=metaconid, entd=entoconid, hypd=hypoconulid.

In MPEF-PV 8166, the preserved portion of the supraspinous fossa is larger than the infraspinous fossa, as in *Anisotemnus* (Shockey and Flynn, 2007). The infraspinous fossa narrows proximally towards the glenoid fossa. A narrow portion of the fossa for the teres minor muscle is present. The spine is high, with a lateral projection equal to the medio-lateral width of the glenoid fossa, and narrower than that of *Anisotemnus* in lateral view (Figure 4.2-3).

In MPEF-PV 8166 the acromion is not preserved. *Thomashuxleya* has a well-developed metacromion with quadrangular shape in lateral view; its presence is uncertain in *Anisotemnus*.

(Shockey and Flynn, 2007). The glenoid fossa is circular, as in *Anisotemnus*, and oriented perpendicular to the main axis of the scapula. In the right scapula of MPEF-PV 8166, the glenoid fossa is taphonomically compressed in the mediolateral plane, resulting in a more oval shape. There is a conspicuous supraglenoid tubercle (Figure 4.1). The coracoid process is well developed and narrower than *Anisotemnus*, although this could be due to diagenetic compression. The length of the coracoid process is smaller than the maximum diameter of the glenoid fossa. The neck is short and wide, similar to *Anisotemnus*.



**FIGURE 4.** Scapulae and humeri of *T. externa* (MPEF-PV 8166). **1** right and **2** left scapulae in proximal view; **3** left scapula in dorsolateral view; right and left humeri in **4** right and left humeri in anterior (top) and distal (bottom) views; **5** right and left humeri in posterior view.

**Humerus.** In MPEF-PV 8166 both left and right humeri are almost complete (Figure 4.4-4.5). The greater tubercle is high, protruding proximally from the level of the head, as in AMNH 28905. The lesser tubercle is smaller than the greater tubercle and much less developed than in *Anisotemnus* (Shockey and Flynn, 2007). The bicipital groove is shallow as in *Anisotemnus* and other specimens of

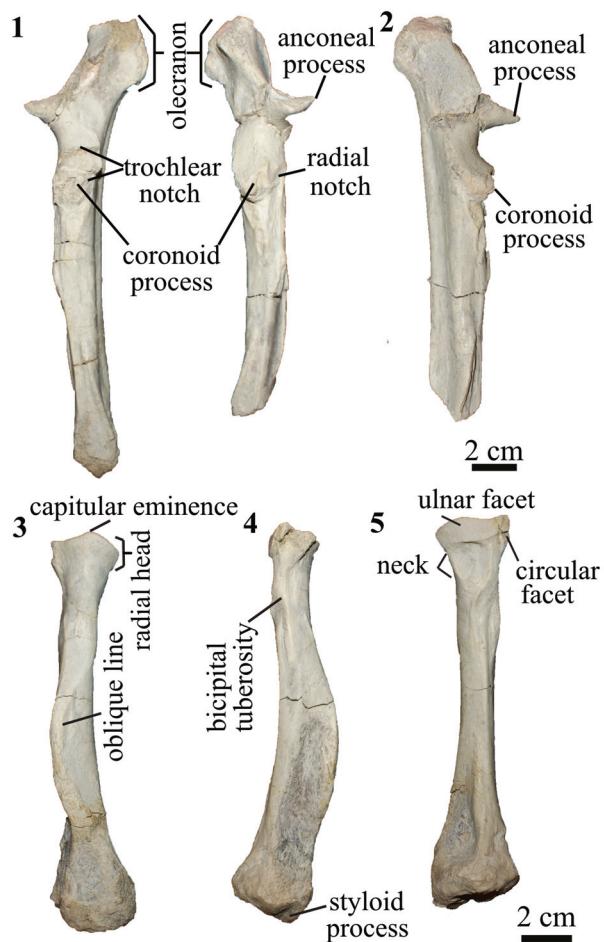
*Thomashuxleya*. The articular surface of the head is oriented posteriorly. The anterior margin of the shaft is convex and the posterior one is straight, and the midshaft is triangular in cross-section. The pectoral (medial) and the deltoid (lateral) crest unite to form a deltopectoral crest, which is large, tilts medially towards its distal end, and extends distal to the midshaft, as in *Anisotemnus* (Shockey

and Flynn, 2007). On the medial side of the shaft, there is a small *Teres major* tuberosity, less developed than that of *Anisotemnus*. The supinator crest is well developed and blade-like as in *Anisotemnus* and *Pleurostyłodon*.

The capitulum is rounded, similar to *Anisotemnus*, and is oblique (i.e., directed proximo-medially) relative to the proximodistal axis, as in other specimens of *Thomashuxleya*, and as opposed to the orthogonal orientation in *Anisotemnus* (Shockey and Flynn, 2007). The medial border of the trochlea is oriented distomedially. The capitulum is wider than the trochlea. The medial crest of the trochlea extends more distally than the capitulum, as in other isotemnids. There is a small, concave capitular tail. The olecranon fossa is shallow (i.e., its depth is less than the maximum diameter of the fossa). The supratrochlear foramen is smaller than the oleocranial fossa. The presence of the supratrochlear foramen in MPEF-PV 8166 can be assessed with confidence because the borders are preserved. Its presence is uncertain in other *Thomashuxleya* specimens (O'Leary et al., 2013), and it is absent in *Pleurostyłodon* and likely absent in *Anisotemnus* (Shockey and Flynn, 2007). The medial condyle is not well preserved, and is not possible to assess the presence of an entepicondylar foramen, which is otherwise present in other *Thomashuxleya* specimens, *Anisotemnus* and *Pleurostyłodon*. The lateral condyle is located more distally than the medial condyle, and there is a distinct radial fossa.

Differences between the humeri of MPEF-PV 8166 and AMNH 28905 (O'Leary et al., 2013) are as follows: in MPEF-PV 8166 the bicipital groove is narrower; the lateral condyle (ectepicondyle) is lateral and not proximal to the capitulum; the coronoid fossa is perforated by the supratrochlear foramen; and the anterior border of the trochlea does not project beyond the plane of the anterior border of the shaft.

**Ulna.** The right and left ulnae are almost complete except for the most distal portion on the left and the incomplete styloid process on the right (Figure 5.1). The shaft is straight. When articulated, the ulna extends distally as long as the radius. Similar to *Anisotemnus* (AMNH 28906), the shaft does not taper distally and is strongly excavated in the lateral side. In *Pleurostyłodon* (AMNH 28904), the lateral margin is less excavated. In lateral view, the shaft is straight along its whole length and not anteriorly concave as observed in other *Thomashuxleya* specimens (e.g., AMNH 28653; Shockey and Flynn, 2007).



**FIGURE 5.** Forelimb of *T. externa* (MPEF-PV 8166). **1** right and left ulnae in frontal view; **2** left ulna in medial view; right radius in **3** frontal, **4** lateral, and **5** posterior views.

The olecranon process of the ulna is long (longer than the trochlear notch) and tilts medially as in *Anisotemnus* and as opposed to *Pleurostyłodon*, where it is straight (Shockey and Flynn, 2007). In lateral view, the olecranon extends posterior to the level of the shaft, in contrast with *Anisotemnus* and *Pleurostyłodon*, where it is more aligned with the shaft (Shockey and Flynn, 2007). The trochlear notch forms a crescent in lateral view and is wider than the midshaft. The coronoid process is as large as the radial notch, in contrast with the larger radial notch in *Anisotemnus* (Shockey and Flynn, 2007). The anconeal process extends laterally and anteriorly (more anterior than the coronoid process), and its width represents about 63% of that of the trochlear notch (Table 4).

The ulnae of MPEF-PV 8166 differ from AMNH 28905 (O'Leary et al., 2013) in the following characters: the absence of a ridge on the lateral

**TABLE 4.** Postcranial measurements of MPEF-PV 8166. \* Translated from Spanish in Elissamburu (2012).

	Value (mm)		Acronym	
	Left	Righth	Scott (1990)	Elissamburu (2012)
Scapula				
Height of the spine	31.0	31.4		
Humerus				
Humerus functional length*	193	204	H1	LFH
Humerus trochlear diameter*	52.0	43.3	H4	DtrH
Maximum length	211	210		
Maximum diameter of the head	44.0	40.5		
Maximum diameter of the mayor tuberosity		46.6		
Maximum diameter of the minor tuberosity		16.6		
Maximum width of the trochlea	30			
Radius				
Maximum length	155	153		
Maximum mediolateral width of the medial shaft	14.1	14.0		
Maximum diameter of the head	32.0			
Minimum diameter of the head	20.3	20.4		
Maximum diameter of the neck	20.3	19.5		
Mediolateral width of the distal epiphysis		36.0		
Diameter perpendicular to the maximum width of the distal epiphysis		31.6		
Ulna				
Anteroposterior diameter of the diaphysis	28.3	28.1		
Olecraneum height	64.9	63.7		
Anteroposterior diameter of the olecranon	32.6			
Proximo-distal length of the trochlear notch	35.9			
Trochlear notch width		32.6		
Width of anconeal process		20.5		
Femur				
Anteroposterior diameter of the shaft	25.5	F7	DAPF	
Mediolateral diameter of the shaft	36.2	F6	DTF	
Tibia				
Mediolateral diameter of proximal epiphysis	55.0	T2	DTpT	
Anteroposterior diameter of proximal epiphysis	44.8			
Lateral condyle transverse width	28.0			
Lateral condyle anteroposterior width	25.0			
Astragalus				
Transverse width of tibial trochlea	27.7		Li 1	
Minimum width of the neck	14.8			
Width sustentacular facet	17.4			
Navicular				
"Dorsal-plantar" width	25.6			
Astragalar facet transverse width	24.0			
Astragalar facet dorsoventral depth	25.0			
Pelvis				
Maximum width of the ilium, above the acetabulum	42.3			

side of the olecranon, the medial inclination of the olecranon, and the absence of a tuberosity on the anterior surface distal to the trochlear notch.

**Radius.** The radius is approximately 70% the length of the ulna (Table 4). The anterior margin of the shaft is convex in lateral view (Figure 5.4) and widens distally, as in *Anisotemnus* and *Pleurostyłodon*. The midshaft is oval in cross-section. The head is oval in proximal view. There is a conspicuous capitular eminence, more developed than in *Pleurostyłodon*, but less than in *Anisotemnus* (Shockey and Flynn, 2007). The bicipital tuberosity is small and placed below an enlarged fossa (Figure 5.4). The ulnar facet is broad and convex (Figure 5.5). There is a circular facet lateral to the ulnar facet as in *Anisotemnus*. It articulates with the radial notch of the ulna, and it does not seem to be a facet for an elbow sesamoid, as observed in *Nesodon* (Scott, 1912) or *Adinotherium* (Croft et al., 2004). The styloid process is short and projects distally, and there is a conspicuous oblique line on the anterior margin (Figure 5.3-5.4). The distal epiphysis is wider than the shaft, and the main axis of the distal articular surface is posterolateral to anteromedial in distal view. In the left radius, the distal epiphysis was diagenetically compressed in the anteroposterior axis.

The radii of MPEF-PV 8166 differ from AMNH 28905 (O'Leary et al., 2013) in having a proximo-lateral radial facet, an oval rather than round mid-shaft cross-section, the presence of an anterior oblique line, and a wide distal epiphysis (but not more than twice as wide as the shaft).

**Pelvis.** Among isotemnidids, the pelvis is only known for *Thomashuxleya* (Simpson, 1936, 1967). In MPEF-PV 8166 the pelvis preserves portions of the right ilium and ischium, but not the pubis (Figure 6.1-6.2). The anterior process of ilium is medio-laterally broad and elongated (Figure 6.1) as in AMNH 28905 (Simpson, 1936; O'Leary et al., 2013); in lateral view (Figure 6.2), the ilium is straight and thin. The acetabulum is circular and oriented posterolaterally. The ischium is in the same anterodorsal plane that the ilium, and projects dorsally from the acetabulum. The posterior part of the ischium is missing.

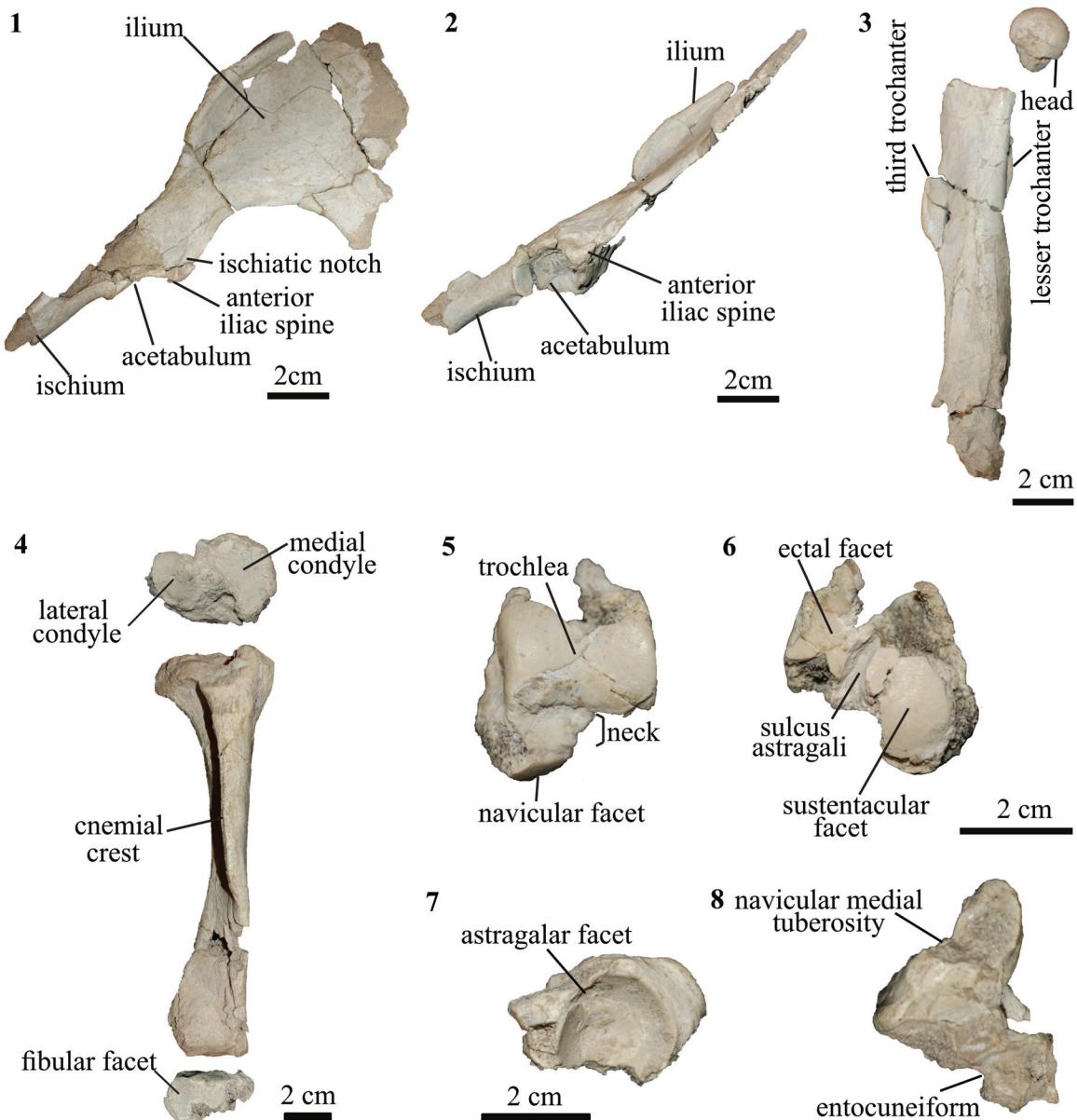
**Femur.** In both femora, the proximal and distal epiphyses are missing and only the shaft and isolated head are preserved (Figure 6.3). Limb suture closure sequence is highly variable in mammals, and several placentals show incomplete fusion of growth plates in adulthood (Geiger et al., 2014). Bone histology from femoral cross-section indicates the specimen was skeletally mature (see

below). This suggests that *Thomashuxleya* had unfused femoral epiphyses into adulthood, as seen in some other mammals (Geiger et al., 2014). The head is spherical with a fovea capitis, enclosed in the articular surface. The trochanteric fossa is deep; the lesser trochanter is smaller than that of AMNH 28905 (O'Leary et al., 2013) and the third trochanter is conspicuous. In lateral view, the anterior border of the shaft is curved and the posterior border is concave.

**Tibia.** In *Thomashuxleya* (MPEF-PV 8166) the tibia and fibula are not fused. In proximal view, the mediolateral width of the proximal epiphysis is greater than the anteroposterior depth, the tibial tuberosity is robust and there are no signs for intercondylar eminences (Figure 6.4). The lateral condyle is circular there is a shallow fossa. The cross section at midshaft is crescent-like, with a lateral concavity. The postero-medial portion of the distal epiphysis of the tibia is not preserved. The distal articulation surface of the tibia with the fibula is small and visible in distal view (Figure 6.4). In other isotemnidids, a partial tibia is only known for an indeterminate isotemnid (AMNH 28690) and *Pleurostyłodon* (AMNH 28904) (Shockey and Flynn, 2007; Simpson, 1936, 1967).

**Astragalus.** The astragalus of MPEF-PV 8166 is the first isotemnid found in association with dental remains that corroborates its species-level identification. It is a left astragalus missing the posterior-most portion of the body (Figure 6.5-6). The astragalus of MPEF-PV 8166 is similar in the overall morphology of AMNH 142463, which, based on its size, was referred to cf. *Thomashuxleya externa* by Shockey and Flynn (2007) (Table 5).

In MPEF-PV 8166 the body is broad and short, as in AMNH 142463. It has a relative shorter neck in comparison with the elements referred to *Thomashuxleya rostrata*, *Pleurostyłodon modicus*, and other isotemnid specimens (Ameghino, 1904b: figures 24, 28-30). The presence of an enlarged medial plantar tuberosity ("medial process" in Shockey and Flynn, 2007; "anterior medial plantar tuberosity" in Szalay, 1994) could not be confirmed as the structure is broken. The trochlear groove is shallow, and the lateral and medial borders of the trochlea are at the same height, in contrast to AMNH 142463 and AMNH 28690 (Isotemnidae indet.) where the lateral border is slightly higher than the medial (Shockey and Flynn, 2007). The trochlea is wider relative to the astragalar length in MPEF-PV 8166 than in AMNH 142463 and the plane of the articulation surface of the lateral facet is almost perpendicular to the trochlear width. The



**FIGURE 6.** Hindlimb of *T. externa* (MPEF-PV 8166). right pelvis in **1** dorsal and **2** lateral views; **3** right femur in anterior view; **4** right tibia in proximal (top), anterior (middle) and distal (bottom) views; left astragalus in **5** dorsal and **6** plantar views; **7** right navicular in proximal view; **8** navicular and entocuneiform in plantar views.

neck is short (Figure 6.5). In dorsal view, the neck represents about half of the trochlear width, similar to AMNH 142463. The navicular facet is convex and anteromedial to the facets for the tibia (Figure 6.5).

The sustentacular facet is elongated antero-posteriorly and covers about half of the astragalus plantar width (Figure 6.6). The proximomedial portion of the sustentacular and the proximal and distal edges of the ectal facets are broken; a well-defined sulcus astragali separates the two. The

ectal facet is restricted to the plantar plane. In ventral view, the medial portion of the navicular facet extends proximally and almost contacts the sustentacular facet (Figure 6.6). There is no cotylar fossa, and the medial astragalar facet is thin and constricted to the medial edge of the trochlea. In MPEF-PV 8166 the most posterior portion of the astragalar body is broken and therefore uninformative regarding the presence of an astragalar foramen, as described for AMNH 142463 and AMNH

**TABLE 5.** Isotemnid astragali measurements in mm.

Specimen No.	Locality	Maximum length	Trochlear width	Maximum width	Head width
MPEF-PV 8166	Cañadón Vaca		27.7		
AMNH 142463	Cañadón Vaca		35.9	41.4	
AMNH FM 14501	"Notostylops beds"	32.8	21.8	32.1	17.7
AMNH 28690	Cañadón Vaca	22	12	20.6	
AMNH 142464	Cañadón Vaca	20.5	8.9	16.7	9.3

28690 (Simpson, 1936, 1967; Shockey and Flynn, 2007).

**Navicular.** MPEF-PV 8166 preserves a right navicular articulated with a fragment of the entocuneiform (Figure 6.7-8). The astragalar facet is circular and deeply concave as described for *Pleurostyłodon* (AMNH 28904) and an indeterminate isotemnid (AMNH 28690; Shockey and Flynn, 2007). The plantar process is present and elongated, with a rounded end. The navicular medial tuberosity is high and conspicuous as in *Colbertia*, *Allalmeia* (Lorente et al., 2014), and AMNH 28690 (Shockey and Flynn, 2007). The cuboid facet is oblique and faces distolaterally, suggesting that the cuboid extended more distally than the navicular. *Thomashuxleya* seems to have the reverse alternate tarsus condition in which the "astragolocuboid contact is lost and calcaneonavicular contact is achieved," following Cifelli (1993, p. 206), although it is not possible to determine if there was a calcaneonavicular contact as the proximal border is broken.

**Metapodials and phalanges.** There is a complete first metapodial, which is long, narrow, and flat. The condyles are asymmetrical, with the lateral one more projected distally than the medial one. The lateral border is concave, and the medial one convex. The proximal epiphysis has a triangular shape and is wide transversally.

Three intermediate phalanges are complete, short, and wide. Two of them are symmetrical, and the remaining phalanx is smaller and asymmetrical. In addition, two ungual phalanges were recovered.

### Body Mass Estimates

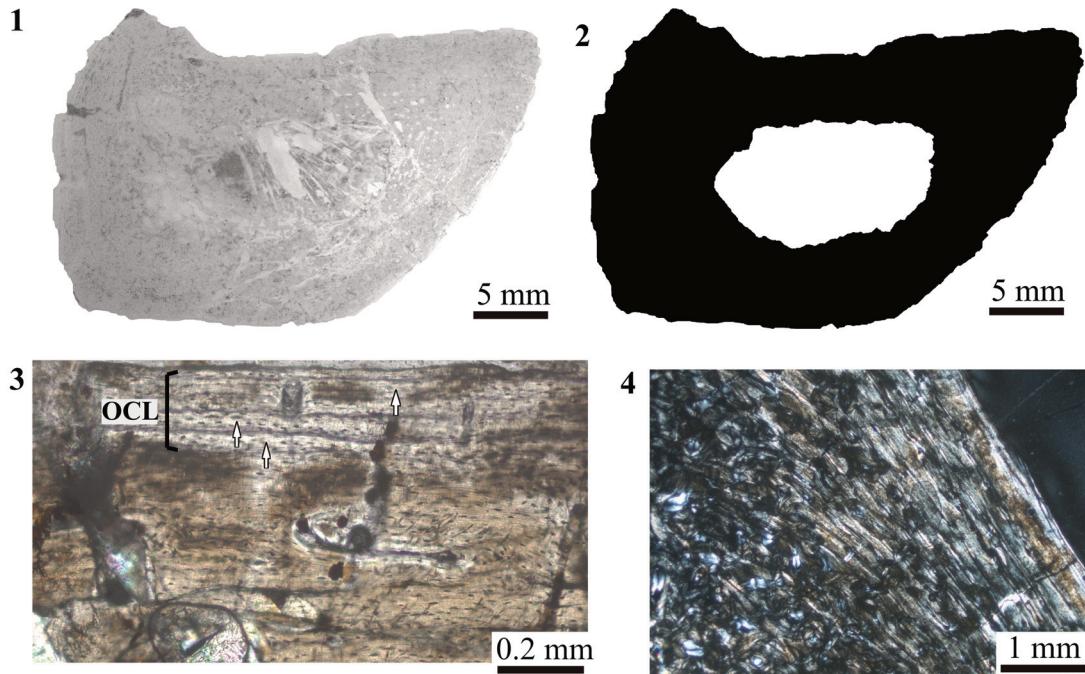
We recovered a wide range of body mass estimates depending on the equation used. The widest range of body mass estimates were obtained with dental measurements (Appendix 1). The second upper molar width gives the highest body mass estimation ( $1501 \pm 583.8$  kg), and the second lower molar length gives the lowest ( $289 \pm 92.2$  kg). Among craniomandibular measurements the posterior jaw length yielded the lowest estimate

( $26 \pm 9.5$  kg), and the maximum width of the mandibular angle yielded the highest ( $209 \pm 84.6$  kg). Regression equations based on postcranial measurements also show considerable variation, but not as much: the highest value is given by the anteroposterior diameter of the femur ( $304 \pm 69.9$  kg) and the lowest by the transverse diameter of the tibia's proximal epiphysis ( $72 \pm 15.1$  kg). The astragalus yielded an estimate of  $84 \pm 24.2$  kg. The arithmetic mean and standard deviation of all the measurements combined were 354.8 kg and 356.2 kg, respectively. The body mass estimations of all the 19 variables analysed have a geometric mean of 235.9 kg and after the bootstrap the 95% confidence interval ranges between 158.1 kg and 354.7 kg. The median is 304 kg, the mode 396 kg, and the kurtosis (a descriptor of the tails of the distribution), is 2.97. As shown in Appendix 1, craniomandibular variables gave a mean of 126 kg and standard deviation of 92.7 kg; dental variables yielded (respectively) 556.4 kg and 390.3 kg; limbs yielded 143.2 kg and 92.9 kg. The estimates using multivariate regression equations yielded a mean of 180.5 kg and a standard deviation of 4.9 kg (Appendix 2).

### Bone Histology

The cross-section shows some sediment in the medullary cavity (Figure 7.1) but can nonetheless be interpreted and represented schematically. *Thomashuxleya* femora display a large open medullary cavity surrounded by the cortex (Figure 7.2), which is not particularly thick in comparison with other large SANUs (Houssaye et al., 2016, figure 3). The image does not allow assessment of the relative thickness of the spongyous transition zone.

Secondary bone of large mammals is typically represented by dense Haversian bone (Kolb et al., 2015b). The bone sample of *Thomashuxleya* is characterized by a compact cortex and a spongyosa medullary cavity (Figure 7.1). The sample shows strong remodelling resulting in dense Haversian bone in the inner cortex (Figure 7.3-4). The primary fibrolamellar bone shows strong laminar



**FIGURE 7.** Bone histology and microstructure of *T. externa* (MPEF-PV 8166). **1** Cross section of the midshaft of the right femur; **2** same as 1 after conversion to a binary image (black represents bone and white the cavities); **3** bone histology under linear polarized light, with black arrows pointing the lines of arrested growth (LAGs). OCL= Outer circumferential layer. **4** Bone histology under cross polarized light.

and longitudinal vascularization. Towards the medullary region, resorption cavities are large and common. An outer circumferential layer (OCL) (Ponton et al., 2004) consisting of avascular lamellar bone is present. The OCL indicates that, despite incompletely fused femoral epiphyses, the specimen reached skeletal maturity (Kolb et al., 2015a; Martinez-Maza et al., 2014). A minimum of three lines of arrested growth (LAGs) could be identified within the OCL (Figure 7.3).

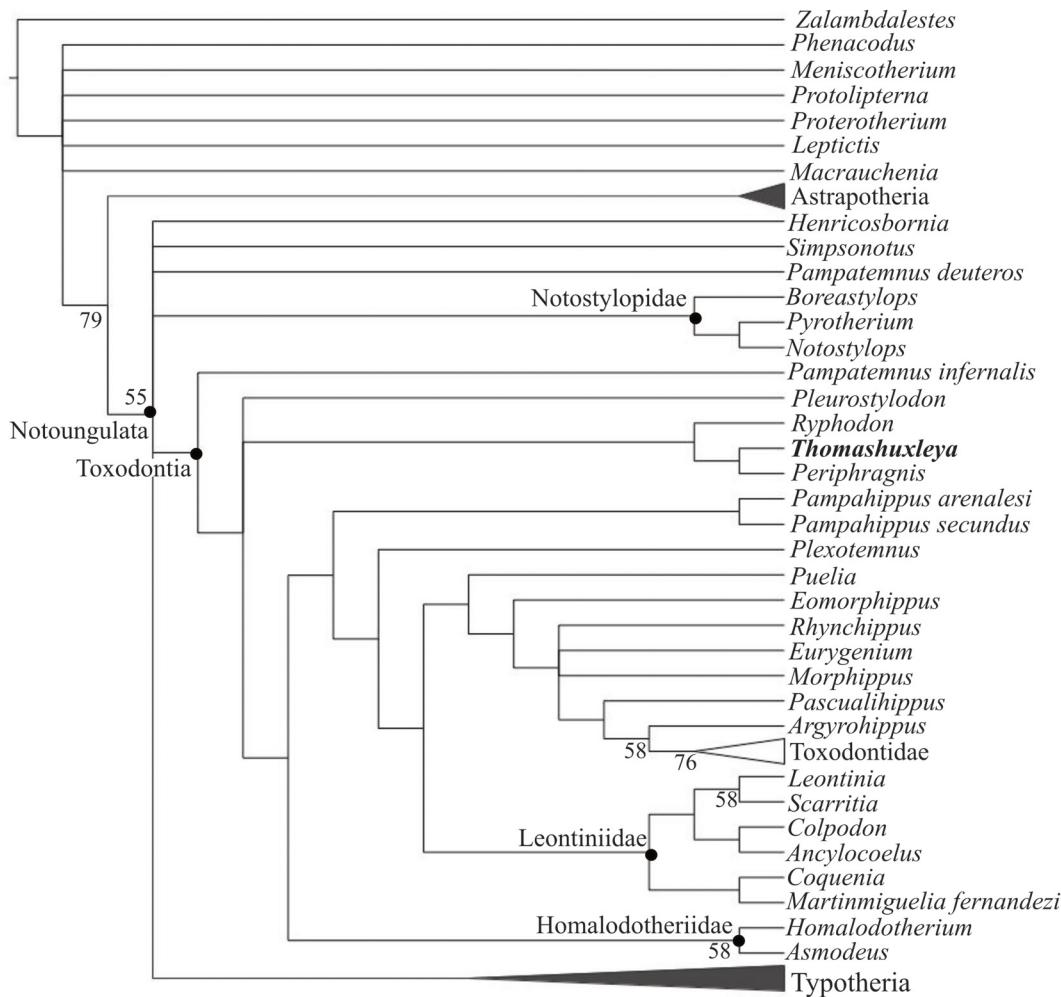
#### Phylogenetic Analysis

**Relationships of *Thomashuxleya* within Notoungulata.** The new *Thomashuxleya* specimen (MPEF-PV 866) provided information to code seven craniodental character states previously unknown for the genus in the character matrix of Deraco and García-López (2015) (see characters [and states] in Morphobank project 2084, 34[1], 35[0], 56[1], 83[0], 96[0], 97[0], 101[0]). The parsimony analysis yielded 281 trees, 420 steps long, with a consistency index (CI) of 0.383 and retention index (RI) of 0.765. A strict consensus of these trees (Figure 8) shows the same topology as that obtained by Deraco and García-López (2015; figure 5), with Notoungulata showing a basal polytomy including *Henricosbornia*, *Simpsonotus*,

*Pampatemnus deuterus*, Notostylopidae (including *Pyrotherium*), Toxodontia and Typotheria. The synapomorphies of the main clades within Notoungulata and Toxodontia are the same as listed by Deraco and García-López (2015; figure 5), and therefore they are not repeated here.

Within Toxodontia, *Pampatemnus infernalis* appears as the most basal taxon. The sister clade shows a polytomy including *Pleurostyloodon*, a (*Ryphodon* [*Thomashuxleya*, *Periphragnis*]) clade, and another consisting of the remaining members of Toxodontia (Figure 8). The synapomorphies supporting the clade (*Ryphodon* [*Thomashuxleya*, *Periphragnis*]) are also the same as listed by Deraco and García-López (2015; figure 5). *Thomashuxleya* has two autapomorphies: the postero-labial fossette on the upper molars disappearing before the closure of the central fossette (34[1]), and the mandibular foramen located at the level of the alveolar border (139[1]).

**Relationships of *Thomashuxleya* within placentalts.** We applied parsimony (MP) using TNT to the combined morphology and protein dataset, including collagens recovered from extant species and Pleistocene fossils (Welker et al., 2015), amino acids for extant taxa (Meredith et al., 2011), and morphology for extinct and extant taxa (O'Leary et

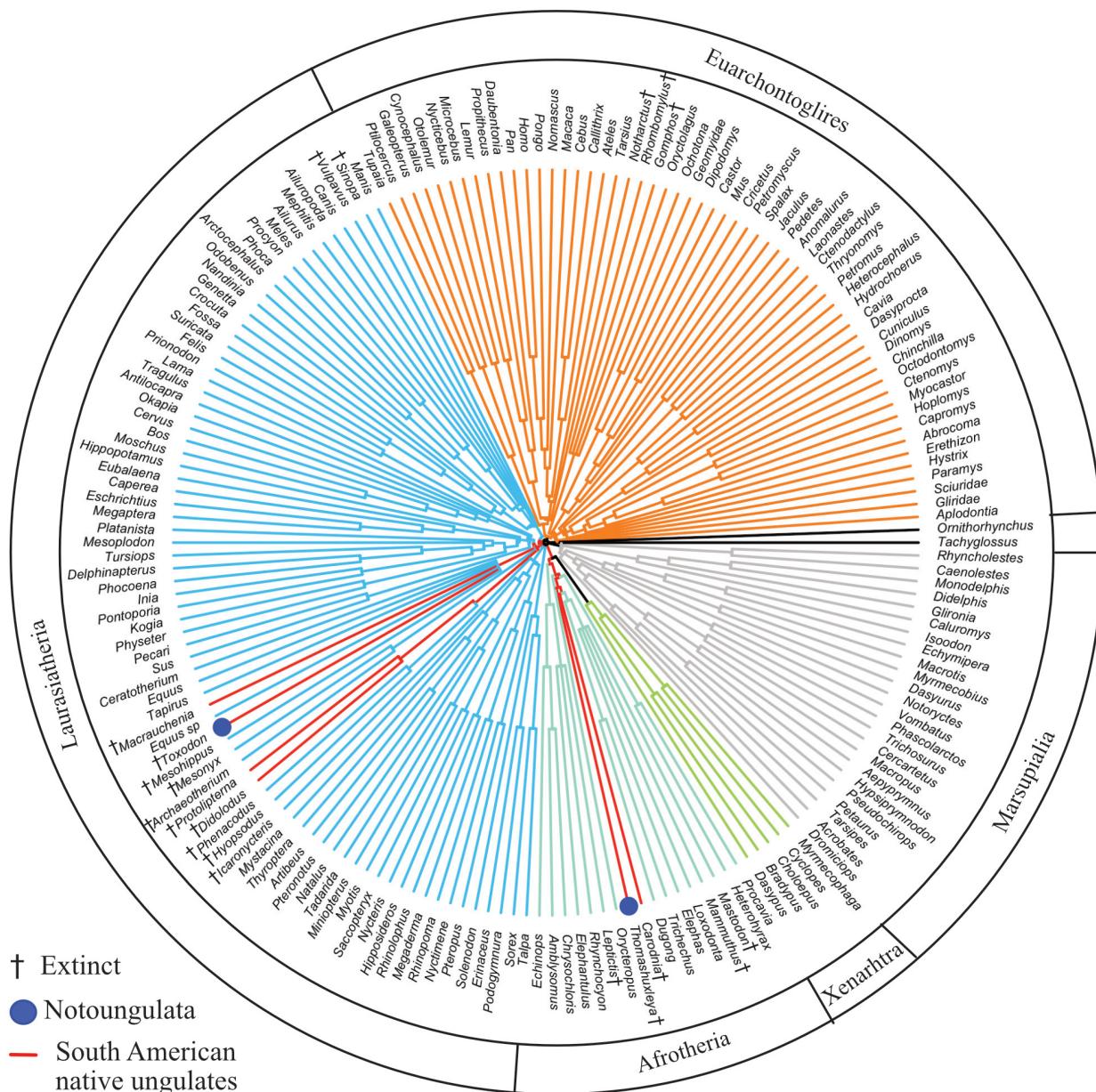


**FIGURE 8.** Strict consensus of 281 trees, 420 steps in length showing the phylogenetic relationships of *Thomashuxleya* within Notoungulata based on the morphological dataset of Deraco and García-López (2015). Numbers indicate bootstrap values above 50.

al., 2013) including our revised anatomical characters for *Thomashuxleya*. This recovered the basic structure of the now well-corroborated tree of placental mammals (Murphy et al., 2001; Tarver et al., 2016). MP analysis with the extended implied weighting yielded 510 trees, 122374 steps long, with a consistency index (CI) of 0.345 and retention index (RI) of 0.638. A strict consensus of these trees (Figure 9) divides meridiungulates between Afrotheria and Laurasiatheria. *Thomashuxleya* and *Carodnia* (sampled for morphology only) are within Afrotheria as sister taxa of tethytheres (i.e., Proboscidea and Sirenia), with a bootstrap value of 10 and 69, respectively. *Toxodon* and *Macrauchenia* (sampled for collagen sequences from Welker et al., 2015) are in a polytomy among perissodactyls, including *Mesohippus* and the fossil *Equus* sp. with

a bootstrap value of 56. *Protolipterna* and *Didolodus* appear in the same clade with *Hyopsodus* and *Phenacodus*, as early divergent members of Euungulata (i.e., perissodactyls and artiodactyls), with a bootstrap value of 88.

We then constrained MP to support the monophyly of each of two clades (but not the two clades together): Notoungulata (*Toxodon* and *Thomashuxleya*) and Litopterna (*Macrauchenia* and *Protolipterna*). This yielded 620 trees of 122391 steps, with a CI of 0.345 and RI of 0.638. In the strict consensus, *Carodnia* is within Afrotheria as sister taxa to tethytheres. The remaining SANUs form a clade with *Hyopsodus* and *Phenacodus*, as sister taxon to Euungulata (Figure 10). The Wilcoxon ranks test shows that there is not a significant difference (p value > 0.05; Table 6) between the Notoungulata



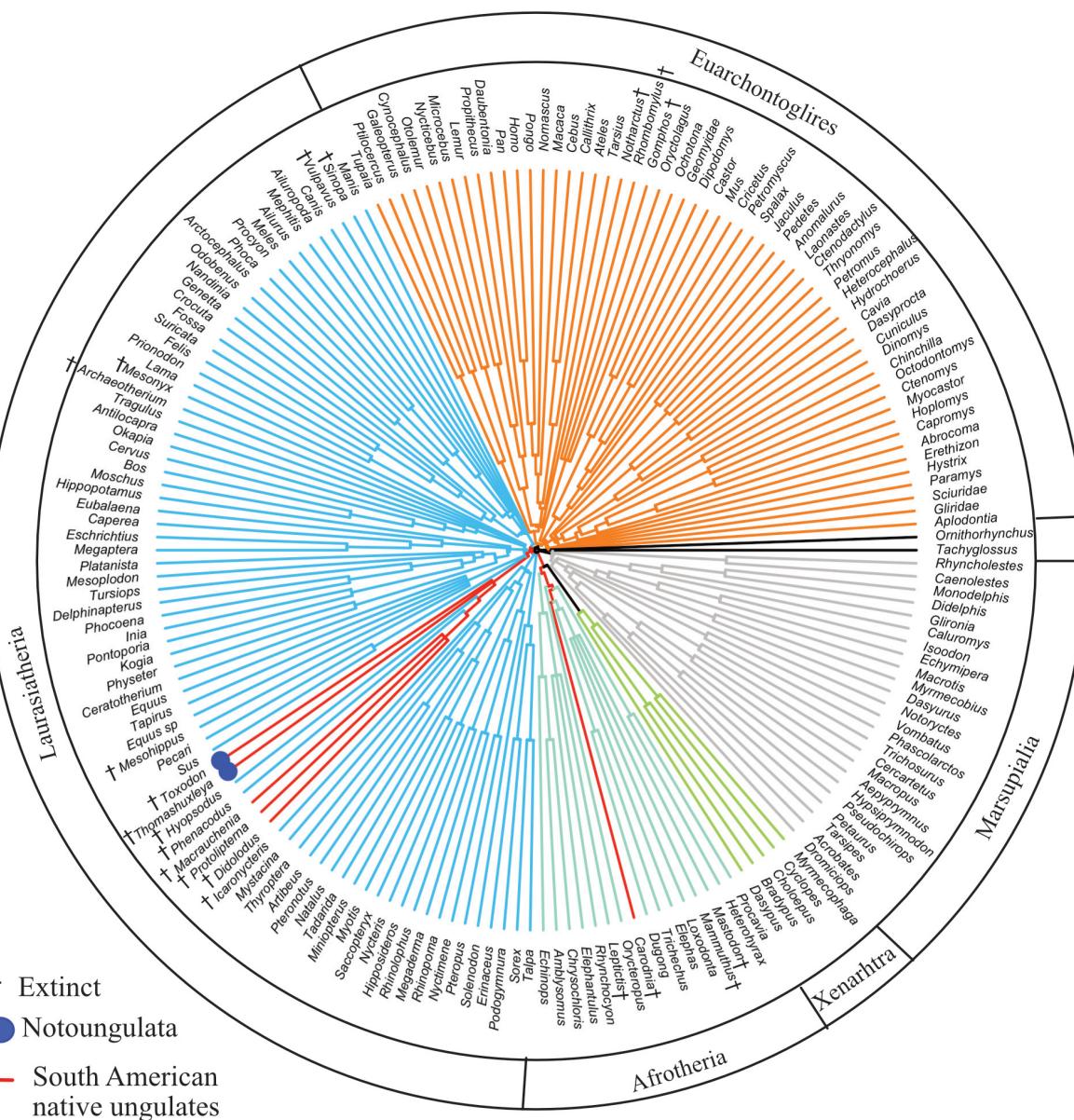
**FIGURE 9.** Strict consensus of 510 trees, 122374 steps in length from unconstrained parsimony analysis of combined proteomic and morphological data.

and Litopterna constraint and the globally optimal topology.

In order to test the different hypothesis regarding the relationships of SANUs, we also constrained all SANUs to form a clade (Meridiungulata). This hypothesis was proposed by McKenna (1975) based on biogeography, but it has not received support by the morphological data (e.g., Horovitz, 2004; Muizon and Cifelli, 2000). The “Meridiungulata” constraint yielded 340 trees of 122424 steps, with a CI of 0.345 and RI 0.638. The strict consensus results in a polytomy with all

SANUs plus *Hyopsodus* and *Phenacodus* forming the sister taxon of Euungulata. Again, a Wilcoxon rank sum test indicates there is not a significant difference ( $p$  value > 0.05; Table 6) between the “Meridiungulata” constraint and the globally optimal topology.

The unconstrained Bayesian analysis yielded an average standard deviation of split frequencies (SDSF) of 0.010 after 3,000,000 generations. This value is well under the recommended SDSF value of 0.05 (Ronquist et al., 2011); however, a number of other metrics identified in Tracer (Rambaut et



**FIGURE 10.** Strict consensus of 620 trees, 122391 steps in length from parsimony analysis of combined proteomic and morphological data constraining monophyly of each of two clades (but not both together): Notoungulata (i.e., *Thomashuxleya* and *Toxodon*) and Litopterna (i.e., *Protopterna* and *Macrauchenia*).

al., 2014) indicate that independent runs did not converge on an optimal topology. The posterior probabilities of the South American fossil taxa with other placentals are low (Figure 11). The unconstrained Bayesian topology shows *Thomashuxleya* as the sister taxon of Xenarthra with a posterior probability of 0.62, *Protopterna* and *Didolodus* as the sister group to Afrotheria (0.88), and *Carodnia* in a clade with *Loxodonta* (0.79). Finally, *Toxodon* and *Macrauchenia* form a clade which appears as

the sister taxon of Perissodactyla with a posterior probability of 0.63.

The same dataset analysed constrained to support Notoungulata (*Thomashuxleya* and *Toxodon*) and Litopterna (*Macrauchenia* and *Protopterna*), as above for MP, yielded an average standard deviation of split frequencies of 0.034 after 3,500,000 generations. The 50% majority rule consensus of the post burn-in topologies (Figure 12) shows Notoungulata and Litopterna as sister clades with a posterior probability of 1. The clade

**TABLE 6.** Results from the Wilcoxon signed ranks test comparing topologies carried out in PAUP. Meridungulata = constraint with all SANUs within a clade; N/L= Notoungulata (*Thomashuxleya*, *Toxodon*) and Litopterna (*Macrauchenia*, *Protolipterna*) monophyly constraints for each clade (but not both together); p values greater than 0.05 indicate no difference between optimal and competing topology.

Topology	Tree length	Rank sums	N	z	p
	122374	(best)			
N/L	122391	52385.0/-48640.0	449	-0.77	0.4390
Meridungulata	122424	214600.5/-193555.5	903	-1.49	0.14

(Notoungulata, Litopterna) appears as sister group of Perissodactyla (0.62) and *Caroindia* is within Afrotheria as sister taxon of *Loxodonta* (0.95) (Figure 12).

## DISCUSSION

### Cranioskeletal Anatomy

The completeness of MPEF-PV 8166 offers a unique opportunity to study the skeletal anatomy of *T. externa* in detail, and provides the first unambiguous associations of cranial and postcranial material from one individual. Dental similarities are substantial among the isotemnids *Thomashuxleya*, *Pleurolystodon*, *Anisotemnus*, and *Periphagnis* (Simpson, 1967), but the dental morphology along with size allowed us to refer MPEF-PV 8166 to *T. externa*. We noted morphological differences in several postcranial elements of MPEF-PV 8166 when compared with other specimens previously referred to this species. In particular, we noted differences in the scapulae, humeri, ulnae, and radii in comparison with AMNH 28905, which was used in O’Leary et al. (2013) to code postcranial characters of this species.

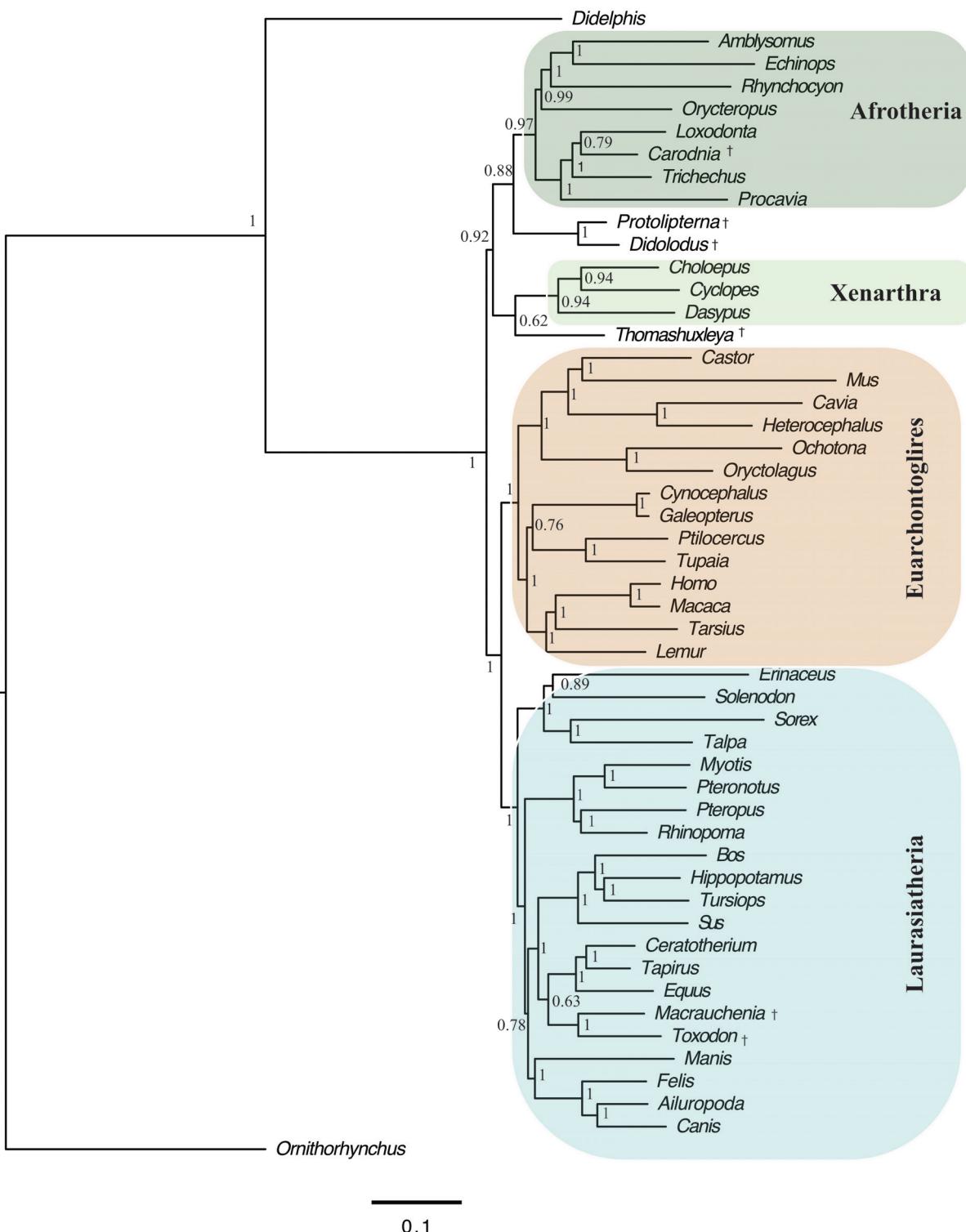
Although incomplete, the astragalus of MPEF-PV 8166 is of particular importance due to its association with craniodental remains. It is slightly smaller than AMNH 142463 referred to *T. externa* by Shockley and Flynn (2007). When compared with other astragali referred to Isotenmidae (Table 5), we found some differences that might indicate species specific characters within this group, such as the relatively high medial and lateral borders of the trochlea, and the relative width of the trochlea in relationship with the astragalar length. *Thomashuxleya* astragali (MPEF-PV 8166 and AMNH 142463) are considerably larger than other specimens variably assigned to this genus (e.g., AMNH 28690, AMNH 142464, and AMNH FM 14501), which therefore probably do not belong to *Thomashuxleya*.

### Body Size

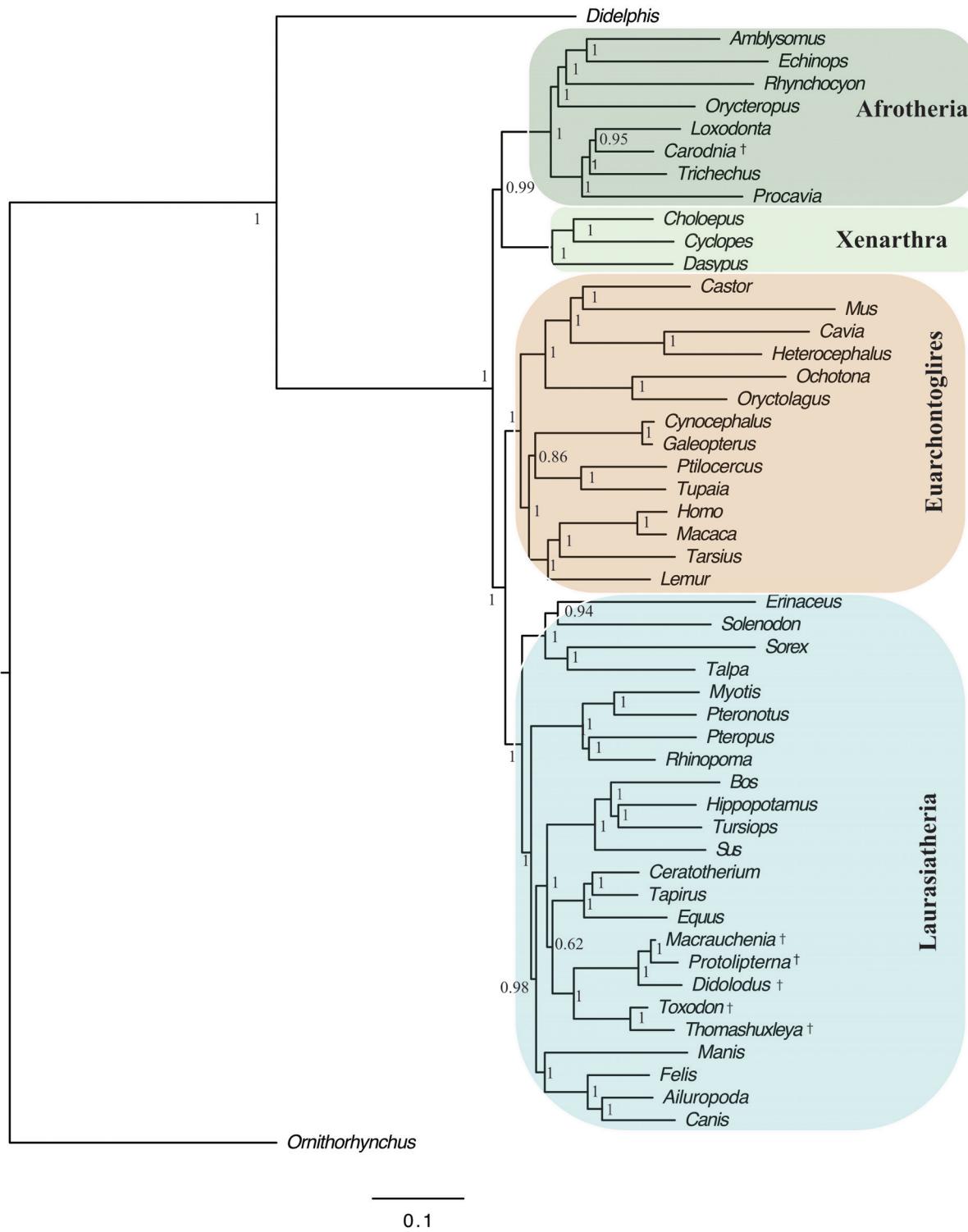
Regression equations obtained from dental variables yielded the highest estimates of body mass of MPEF-PV 8166 (Appendix 1), with an arithmetic mean of 556.4 kg. Other subsets of measurements differed in their estimates: crano-mandibular variables gave a mean of 126 kg, limbs yielded a mean of 143.2 kg, multivariate regression equations that included dental variables yielded a mean of 180.5 kg (Appendix 2), and the body mass estimate of the astragalus was  $84 \pm 24.2$  kg.

For Isotenmidae, Elissamburu (2012) favored estimations obtained from dental measurements. The dental dimensions can be influenced by differences in function and diet, which challenges their use to estimate the body mass in fossil mammals with no living descendants. Damuth (1990) noted that Paleogene ungulates from North America tend to have larger teeth relative to their body mass than extant ungulates. Considering the values obtained from the postcranial bones, we believe dental dimensions overestimate the body mass of MPEF-PV 8166 (Appendix 1).

Proximal limb bones are good estimators of body mass because they are weight-bearing elements, subject to biomechanical constraints (Scott, 1990). Elissamburu (2012) considered some postcranial measurements to be good estimators of body mass for *Thomashuxleya*, but not for closely related taxa such as *Pleurostylodon*, which is odd given the comparable dimensions and anatomy of both taxa. MPEF-PV 8166 and other specimens referred to *Thomashuxleya* show robust limbs. MPEF-PV 8166 exhibits some morphological features which are suggestive of scratch-digging capabilities, such as a high spine of the scapula, a large humeral deltopectoral crest that extends distally, and a long ulnar olecranon (Shockley et al., 2007). Fossilized habits have been inferred in mesotheriids (Shockley et al., 2007), and other notoungulates (e.g., *Protypotherium*) show features that suggest digging capabilities (Croft and Anderson, 2008). Fossilized habits may be an ancestral attribute in notoungulates (Shockley et



**FIGURE 11.** Optimal Bayesian tree (i.e., 50% majority rule of post-burn-in trees) of combined proteomic and morphological data. Numbers represent Bayesian posterior probabilities; daggers indicate fossil taxa.



**FIGURE 12.** Optimal Bayesian tree (i.e., 50% majority rule of post-burn-in trees) of combined proteomic and morphological analysis constraining monophyly of each of two clades (but not both together): Notoungulata (i.e., *Thomashuxleya* and *Toxodon*) and Litopterna (i.e., *Protolipterna* and *Macrauchenia*). Numbers represent Bayesian posterior probabilities; daggers indicate fossil taxa.

al., 2007; Croft and Anderson, 2008), and if this is the case, limb bones could also overestimate the body size in notoungulates. The estimates obtained from postcranial measurements (mean 143.2 kg) are higher than the estimate of 70–90 kg for *Thomashuxleya* of Croft (2016). This estimate was based on a value of ca. 1.5 m of head-body length based on Simpson's skeletal reconstruction (Croft, personal commun., 2017). In extant ungulates, the regression of head-body length shows the highest correlation with body mass and lowest percent of prediction error (Damuth, 1990).

The body mass estimate from the astragalus ( $84 \pm 24.2$  kg) is congruent with the estimate of 70–90 kg of Croft (2016) based on the estimated head-body length of *Thomashuxleya*. Astragalar regression equations of Tsubamoto (2014) are based on a broad taxonomic sample of mammals, whereas the regression equations using limb bones presented by Scott (1990) are based on living ungulates (artiodactyls and perissodactyls). Of the variety of body mass estimates for MPEF-PV 8166, we favor the one from the astragalus as it is based on a broader taxonomic sample of living mammals, and taken into account that morphological features in the limb bones associated with scratch-digging capabilities might bias the body mass estimate of MPEF-PV 8166.

*T. externa* is one of the largest mammalian taxa of the Vacan (Cifelli, 1985), only comparable in size with the astrapothere *Albertogaudrya*, according to the length of the lower molars. *T. externa* is the earliest known, anatomically well-documented toodoxodontian notoungulate (Billet, 2011; Deraco and García-López, 2015). Basal notoungulates such as *Henricosbornia*, which also occur in the Vacan but with an origin in at least the Itaboraian (early Eocene; ~53–50 Ma; Gelfo et al., 2009; Woodburne et al., 2014a), were much smaller. For example, the m<sub>2</sub> length of *Henricosbornia waitehor* from the Rio Chico Formation (Simpson, 1935) comprises less than a quarter of the m<sub>2</sub> length in MPEF-PV 8166. With an estimated body mass of ~84 kg, *Thomashuxleya* demonstrates a large range of notoungulate body masses by the middle Eocene, and its postcranial morphology can be expected to reflect weight-bearing adaptations.

To our knowledge, of the more than 150 species recognized for notoungulates, bone histology is documented in only four (de Ricqlès et al., 2009; Tomassini et al., 2014, 2015, 2017; Forasiepi et al., 2015; Kolb et al., 2015b). As in *Thomashuxleya*, the toxodontians *Toxodon* and *Nesodon* showed

Haversian bone with a compact cortex, a feature associated with the increased loading in large-bodied mammals (Straehl et al., 2013; Forasiepi et al., 2015; Kolb et al., 2015b; Tomassini et al., 2015). The presence of an outer circumferential layer (OCL) in the histological section of MPEF-PV 8166 indicates the specimen represents a skeletally mature individual of *T. externa*. At least some wear is evident on all of its dental loci, although compared to other Vacan specimens (and indeed Paleogene SANUs in general; see Strömberg et al., 2013) the degree of wear is fairly low.

Houssaye et al. (2016) analysed the bone microstructure of several SANUs including the toxodontid *Nesodon*. The femora of this taxon showed a large open medullary cavity surrounded by a spongy zone. A qualitative and quantitative analysis of the bone microstructure strongly suggest graviportal for *Nesodon* (Houssaye et al., 2016), based on the relatively high bone compactness and cortex thickness, and its position in a bone microstructure morphospace close to graviportal living mammals such as the rhinos *Ceratherium* and *Dicerorhinus*.

## Systematics

The craniodental anatomy of MPEF-PV 8166 is similar to other specimens previously referred to *Thomashuxleya*. The completeness of the new *Thomashuxleya* specimen enable us to revise and add to data known for this taxon in a phylogenetic analysis within Notoungulata and within Placentalia. The craniodental anatomy of MPEF-PV 8166 supports the placement of *Thomashuxleya* as an early divergent taxon within Toxodontia (Billet, 2011; Deraco and García-López, 2015). Some alleged synapomorphies that have previously been used to link SANUs with Afrotheria (Agnolin and Chimento, 2011) have already been convincingly refuted (Billet and Martin, 2011; Kramarz and Bond, 2014). Other studies group early members of Litopterna and South American “condylarths” with certain North American “condylarths” (Muizon and Cifelli, 2000; Gelfo, 2007) apparently a part of Laurasiatheria (Halliday et al., 2017). O’Leary et al. (2013) placed *Thomashuxleya* and *Carodnia* within Afrotheria, and *Didolodus* and *Protolipterna* within Laurasiatheria. Halliday et al. (2017) included *Protolipterna* and the notoungulate *Simpsonotus* in a comprehensive phylogenetic study of Paleocene placentals. In their study, *Protolipterna* was closely related to archaic artiodactyls, but the position of *Simpsonotus* was inconsistent, varying among different placental clades.

Welker et al. (2015) presented a new source of data to address this question in a pioneering study that used collagen protein sequences from two Pleistocene species to test the phylogenetic affinities of meridiungulates. They hypothesized that at least some of South America's endemic mammalian groups are close relatives of Perissodactyla. Previous authors (e.g., Billet, 2011; Cifelli, 1993; Gelfo, 2007; Horovitz, 2004; Muizon and Cifelli, 2000; Simpson, 1936, 1967) have shown that the two most speciose groups (notoungulates and litopterns) are ungulate-grade placentals; thanks to the novel data from Welker et al. (2015), we can now attempt a greater level of precision.

To fully evaluate the impact of the new material (if any) on the estimation of the relationships of *Thomashuxleya*, at least some integration of biomolecular and gross anatomical data is necessary. The matrix of O'Leary et al. (2013) is, to date, the largest available anatomical matrix that also samples at least some meridiungulates, and we have improved on their coding based on the new material described here. Further progress will be gained once the comprehensive matrix of O'Leary et al., 2013 is revised in both definitions and character codings; our study has made a start on the latter. The multiple analyses we perform is also a case study in which different data sources are integrated and results are evaluated from multiple perspectives. We believe that the combination of morphological and molecular data can provide more precise and robust hypotheses for several extinct clades of mammals (e.g., Asher et al., 2005; Muizon et al., 2015; Pattinson et al., 2015), and we further note that combination of molecular and morphological datasets can in principle increase support for clades not present when one or both are analysed in isolation (Gatesy and Baker, 2005; Lee and Camens, 2009; Thompson et al., 2012).

On the other hand, we do not maintain that combination of phylogenetic data from multiple sources (e.g., collagens from recent taxa, morphology from ancient fossils) will in every case yield a well-supported topology. The variety of optimality criteria we have applied to our analysis of phylogenetic data including meridiungulates and modern taxa does not confirm either the relationship proposed by Welker et al. (2015) with Perissodactyla, nor does it rule out alternatives that place one or more SANUs close to Afrotheria, Euungulata, or even Xenarthra. The strict consensus of our most parsimonious trees split Notoungulata, placing *Thomashuxleya* within Afrotheria and *Toxodon* within Laurasiatheria (Figure 9). The post-burnin,

majority rule consensus of our optimal Bayesian trees place *Thomashuxleya* as the sister taxa of Xenarthra (with a very low posterior probability) and *Didolodus*, *Protolipterna*, and *Carodnia* in or near Afrotheria (also with low posterior probabilities; Figure 11). The monophyly of Notoungulata and Litopterna is supported by morphological data of the extensive fossil record of these two groups. The MP and Bayesian analyses constrained to support the monophyly of Notoungulata (*Toxodon* and *Thomashuxleya*) and Litopterna (*Macrauchenia* and *Protolipterna*) recovered these clades within Laurasiatheria (Figure 10 and 12), a result that approximates the hypothesis of Welker et al. (2015) that one or more meridiungulate groups are closely related to Perissodactyla.

There are a number of causes potentially behind this lack of a confident resolution of the phylogenetic relationships of meridiungulates with other placentals. First, although the new *Thomashuxleya* specimen is one of the oldest and most complete notoungulate yet known, it is still a highly derived taxon, and its potential synapomorphies with any extant relatives have been at least partly overwritten by homoplasy. This possibility is demonstrably not the case for many vertebrate groups with an extensive Paleogene fossil record (e.g., primates as shown by Pattinson et al., 2015), but it is naive to expect that such a scenario has never happened during the course of mammalian evolution. Muizon et al. (2015) performed a phylogenetic analysis of placental interordinal relationships using morphological characters and including SANUs and a large sample of other Paleogene fossils. They pointed out the high level of homoplasy detected in their dataset, and our data also clearly demonstrate homoplasy (RI of 0.638 and CI of 0.345 in our unconstrained tree).

Second, some part of the anatomy, which we have not yet sampled (e.g., inner ear), may yet contain key phylogenetic information with the potential to resolve the ambiguity discussed here (Macrini et al., 2010, 2013). An ear bone referred to an early divergent notoungulate shows that this region exhibits a mixture of plesiomorphic and derived features for Notoungulata (Billet and Muizon, 2013). The ear region of notoungulates exhibits similarities with hyracoids (Billet and Muizon, 2013), but further investigation is required to determine their phylogenetic implications. Finally, and non-exclusively, the early radiation of Notoungulata and other SANUs occurred very rapidly on parts of the continent which are not yet well sampled. Early Paleogene localities (e.g., Tiupampa, Itaborai)

older than Cañadón Vaca exist, but have not yet yielded well-preserved, associated remains of notoungulates. Although it is a Xenungulate and not a notoungulate, *Carodnia* is a notable exception, comprising another large, Eocene meridiungulate known from associated cranial and postcranial remains from Itaborai, Brazil. Accordingly, phylogenetic uncertainty may dissipate with the inclusion of additional anatomical characters (e.g., those derived from CT imaging) and with the discovery of more complete notoungulate material predating the Eocene. Further extraction of durable protein sequences from other meridiungulates, even those that predate the Neogene, remains theoretically possible and is clearly among the most promising avenues for further research into meridiungulate systematics (cf., Welker et al., 2015).

## CONCLUSIONS

MPEF-PV 8166 is one of the oldest, most complete associated notoungulate skeletons yet documented in the literature. We refer this specimen to *T. externa* based on dental morphology and size. The presence of an associated dentition with postcrania enables us to recognize variation in skeletal elements previously assigned to this taxon. *T. externa* weighed approximately  $84 \pm 24.2$  kg. Dental measurements provided larger estimates for body mass than skeletal elements. *T. externa* demonstrates that notoungulates acquired large size by the middle Eocene, and its bone histology shows that MPEF-PV 8166 was a skeletally mature individual with the capacity to withstand heavy loading, despite its incompletely fused femoral epiphyses. Our phylogenetic analysis raises a number of possibilities regarding its affinities. With *Thomashuxleya* constrained to be part of a monophyletic Notoungulata, MP recovers *Thomashuxleya* near the base of euungulates, close (but not on the stem) to Perissodactyla, whereas Bayesian analysis places it on the stem to Perissodactyla. Unconstrained MP and a Bayesian analysis splits meridiungulates into two groups: Pleistocene taxa known for collagen proteins with euungulates, and Paleogene taxa with afrotherians and xenarthrans.

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**APPENDIX 1.**

Definitions, measurements (in mm, unless otherwise stated) and equations used (Janis, 1990; Scott, 1990; Tsubamoto, 2014) to estimated the body mass (BM) in kg of MPEF-PV 8166. PE= Percent of error.

Variable	Acronym	Definition	Type	Dataset	Value	Equation	R <sup>2</sup>	% PE	Slope	Intercept	BM	BM + BM- PE	
Lower molar row length	LMRL	Measured along the base of the teeth	Dental	All ungulates	75.6	log BM = 3.265*log(LMRL)- 0.94 0.536	31.9	3.27	-0.54	396	522	269	
First lower molar length	FLML	Measured at the occlusal surface of the tooth	Dental	All ungulates	20.1	log BM = 3.263*log(FLML)+1.337	0.93	34.6	3.26	1.34	389	524	254
First lower molar width	FLMW	FLMA= FLML*FLMW	Dental	All ungulates	15.6	log BM = 2.909*log(FLMW)+2.030	0.92	38.4	2.91	2.03	316	437	194
First lower molar area	FLMA	FLMA= FLML*FLMW	Dental	All ungulates	313.3	log BM = 1.553*log(FLMA)+1.701	0.93	33.2	1.55	1.7	378	503	252
Second lower molar length	SLML	Measured at the occlusal surface of the tooth	Dental	All ungulates	22.5	log BM = 3.201*log(SLML)+1.130	0.94	31.9	3.2	1.13	289	381	197
Second lower molar area	SLMA	SLMA= SLML *Second Lower Molar Width	Dental	All ungulates	343.7	log BM = 1.563*log(SLMA)+1.541	0.94	33.5	1.56	1.54	320	427	213
Third lower molar area	TLMA	TLMA= Third lower molar length *Third lower molar width	Dental	All ungulates	508.9	log BM = 1.580*log(TLMA)+1.404	0.93	33.1	1.58	1.4	479	638	321
Second upper molar length	SUML	Measured at the occlusal surface of the tooth	Dental	All ungulates	27.9	log BM = 3.184*log(SUML)+1.091	0.93	34.7	3.18	1.09	494	666	323
Second upper molar width	SUMW	Measured at the occlusal surface of the tooth	Dental	All ungulates	36.9	log BM = 3.004*log(SUMW)+1.469	0.92	38.9	3	1.47	1501	2085	917
Second upper molar area	SUMA	SUMA=SUML*SUMW	Dental	All ungulates	1029.5	log BM=1.568*log(SUMA)+1.27 7	0.94	32.7	1.57	1.28	1002	1329	674
Posterior jaw length	PJL	Measured as the horizontal distance from the junction of the posterior part of m3 with the jaw to the posterior border of m <sub>3</sub>	Crania	All ungulates	65.3	log BM= 2.412*log(PJL)+0.031 0.352	0.93	36.5	2.41	0.03	26	35	16
Maximum width of the mandibular angle	WMA	Measured from the junction of the posterior part of m3 with the jaw to the maximally distant point on the angle of the jaw	Crania	All ungulates	105.5	log BM= 2.803*log(WMA)- 0.92 0.352	40.5	2.8	-0.35	209	293	124	
Length of the ridge for the masseteric attachment	MFL	Measured from the posterior portion of the jaw glenoid to the most anterior extent of the scar for the origin of the masseter muscle	Crania	All ungulates	153	log BM= 2.950*log(MFL)- 0.94 1.289	35	2.95	-1.29	143	193	93	
Humerus functional length (cm)	H1	The distance between the most proximal point of the head and the most distal point of the troclea	Postcranial	All ungulates	19.85	log BM= 3.4026*log(H1)- 0.92 2.3707	28	3.4	-2.37	111	142	80	
Humerus trochlear diameter (cm)	H4	The transverse distance of the distal articular surface	Postcranial	All ungulates	4.76	log BM= 2.4815*log(H4)+0.4516 0.95	22	2.48	0.45	136	166	106	

Variable	Acronym	Definition	Type	Dataset	Value	Equation	R <sup>2</sup>	% PE	Slope	Intercept	BM	BM + PE	BM - PE
Anteroposterior diameter of the femur (cm)	F6	The transverse diameter of the diaphysis at its midpoint	Postcranial	All ungulates	3.62	BM=2.82*10 <sup>log(F6)+0.9062</sup>	0.94	23	2.82	0.91	304	373	234
Mediolateral diameter of the shaft (cm)	F7	The anteroposterior diameter of the diaphysis at its midpoint	Postcranial	All ungulates	2.55	log BM=2.6016*log(F7)+0.9119	0.94	26	2.6	0.91	93	117	69
Diametro transverso proximal de la tibia (cm)	T2	The transverse diameter of the tibia's proximal epiphysis	Postcranial	All ungulates	5.5	log BM=2.8491*log(T2)-0.2495	0.95	21	2.85	-0.25	72	88	57
Transverse width of tibial trochlea	L1		Astragalus	Various mammals	27.7	In BM= 2.789*ln(L1)+2.078	0.98	28.83	2.79	2.08	84	109	60

**APPENDIX 2.**

Selected measurement (in cm) and multivariate equations used (Mendoza et al., 2006) to estimated the body mass in kg (BM) of MPEF-PV 8166.

Adj. R <sup>2</sup>	%MPE	mid PE	LMRL	LPRL	JMA	JMC	JD	JMB	BM+ mid		BM -mid	
									BM	PE	BM	PE
0.98	21-25	23	7.56	5.86	6.53	10.55	4.88		177	217	167	
0.98	21-25	23	7.56	5.86	6.53	10.55	4.88	11.6	184	226	174	