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Adaptive Radiation of Multituberculate Mammals Before the Extinction of Dinosaurs

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1. Dental complexity data collection

We scanned lower cheek tooth rows (premolars and molars) of 41 multituberculate genera using a Nextec Hawk three-dimensional (3D) laser scanner at between 10 and 50- μ m resolution and in a few cases a Skyscan 1076 micro X-ray computed tomography (CT) at 9- μ m resolution and an Alicona Infinite Focus optical microscope at 5- μ m resolution. One additional scan from the PaleoView3D database (<http://paleoview3d.marshall.edu/>) was made on a Laser Design Surveyor RE-810 laser line 3D scanner with a RPS-120 laser probe. 3D scans of cheek tooth rows are deposited in the MorphoBrowser database (<http://morphobrowser.biocenter.helsinki.fi/>). The

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sampled taxa provide broad morphological, phylogenetic, and temporal coverage of the Multituberculata, representing 72% of all multituberculate genera known by lower tooth rows. We focused at the level of genera rather than species under the assumption that intrageneric variation among multituberculates is mostly based on size not gross dental morphology. The sampled cheek tooth rows for each genus are composed of either individual fossil specimens or epoxy casts with complete cheek tooth rows or cheek tooth row composites formed from multiple fossil specimens or epoxy casts of the same species. *Barbatodon* and *Hainina* are exceptions, in which composites were constructed from specimens representing multiple species of each genus. For some genera, more than one cheek tooth row was scanned, representing either multiple individuals of a single species or multiple species from the same genus.

Scan data were digitized, processed, and analyzed as in Evans et al.¹⁸. The 3D point files were converted into digital elevation models (DEMs) of the cheek tooth rows. To focus on shape apart from size, all cheek tooth rows were scaled to a length of 150 pixel rows. The number of ‘tools’ on the crown or the ‘dental complexity’ was approximated by GIS analyses of the full 3D shape of the occlusal surface of the cheek teeth. The specific measure of ‘dental complexity’ that we used is ‘orientation patch count’ (OPC). It is calculated by first determining the surface orientation at each pixel on the DEM. Then contiguous pixels that are facing the same cardinal direction (e.g., north, south, southwest) are grouped into patches. The number of these ‘patches’ is the ‘orientation patch count’ or the OPC. To reduce the effect of a specific tooth orientation, the OPC calculation was repeated eight times at rotations of multiples of 5.625° and the mean of these

repetitions is used (OPCR). For the dataset of Evans et al.¹⁸, OPC and OPCR are highly correlated (adjusted $R^2 > 0.98$ for both upper and lower), with the average deviation of OPC from OPCR being 3.00%. The same analyses in Evans et al.¹⁸ were re-run on the OPCR data and no differences were found in recorded patterns (Supplementary Table 6).

Although degree of wear among specimens varies and despite studies having shown that specific functional features may be modified by tooth wear³⁰, general topographic measures tend to be more stable. This suggests that higher-level patterns, like OPC, are relatively robust to wear³¹. Specifically in the case of OPC, all slopes irrespective of their steepness are included in calculations; hence even worn cusp features are typically tabulated. Nevertheless, we excluded fossil specimens with excessive tooth wear or post-mortem abrasion. Specimens with moderate/heavy wear that were included in the analysis are in boldface in Supplementary Table 1. When we excluded those specimens with moderate/heavy wear from the analyses, the pattern of results remained the same. Note that the use of worn specimens typically underestimates OPC and thus decreases the amount of herbivory inferred.

To examine intraspecific variation of OPC, the coefficient of variation (SD/Mean) for 18 specimens of *Apodemus flavicollis* unworn lower tooth rows was 8.56% (mean \pm SD: 216.85 ± 18.56). The relative standard error ($(SD/\sqrt{n})/\text{Mean}$) is 2%.

To test for the compatibility of data from different scanning systems, two multituberculate tooth row casts (*Meniscoessus robustus* UCMP 107405, *Parikimys carpenteri* DMNH 52224) were scanned with a Nextec Hawk point laser scanner (University of Helsinki), Skyscan 1076 MicroCT scanner (University of Washington,

Seattle), and a Laser Design Surveyor DS2025 laser line scanner (Monash University, Melbourne) at 10 μm resolution (Supplementary Figures 9 and 10). The relative standard error of the measurements of OPC values for the tooth rows, downsampled to 150 data rows, was less than 3.1% (Supplementary Table 7). We were unable to include the Alicona Infinite Focus optical microscope in this comparison. Nevertheless, only two Late Jurassic specimens were scanned using this machine. When they are removed from the dataset, there is no effect on the major results of the study.

For genera with more than one sampled cheek tooth row, OPC values varied by no more than 15% of the mean OPC. For these genera, we used the mean OPC value. The specimens used in each of our sampled cheek tooth rows and their OPC values are listed in Supplementary Table 1 and plotted in Fig. 2a as solid horizontal lines, lengths of which correspond to the temporal range of the genus or uncertainties in the age of the fossil localities. Resolution of temporal range data varies from geologic ages to land-mammal ages constrained by radiometric ages. We gratefully acknowledge the museums and institutions listed in Supplementary Table 1, and we would also like to thank the individuals that made these specimens, casts, and scans available: Zofia Kielan-Jaworowska, Dave Krause, Gregg Gunnell, Philip Gingerich, Toni Culver, Jaelyn Eberle, Logan Ivy, William A. Clemens, Patricia Holroyd, Matthew Carrano, Michael Brett-Surman, Nicholas D. Pyenson, Richard C. Fox, Rich Cifelli, Brian Davis, Meng Jin, Ivy Rutzky, Douglas Boyer, Ruth O'Leary, Zoltan Csiki, Thomas Martin, Aisling Farrell, Luis Chiappe, Guillermo Rougier, Suzanne Strait, Jack Horner, Phillipa Brewer, and Jerry Hooker; Bryan Small for cast preparation; William Sanders for specimen

preparation; and Suzanne Strait, Douglas Boyer, Norman Macleod, Jonathan Krieger, Murat Maga, Casey Self, Abby Vander Linden, and Timothy Cox for help with scan data.

To compensate for the relatively poor fossil record from the Late Jurassic and Early Cretaceous (155–100 Myr ago), we developed formulae to estimate OPC values that are based on the positive correlation of OPC and the proportional length of the lower molars in the cheek tooth row ($[\text{length of } m1 + m2]:[\text{length of all lower premolars}]$) of sampled multituberculates (Supplementary Fig. 1; $r = 0.750$, $p < 0.0001$, $n = 40$) and the ratio of the lower first molar to the lower fourth premolar ($\text{length of } m1:\text{length of } p4$) (Supplementary Fig. 2; $r = 0.711$, $p < 0.0001$, $n = 40$). The resulting estimated OPC values for Late Jurassic and Early Cretaceous taxa (dashed lines in Fig. 2a; Supplementary Table 2) fall within or close to the range of measured OPC values for this interval, suggesting that the pattern for this interval is robust to gaps in the sampling. In Fig. 2a, mean and standard deviation of OPC were calculated for 5-Myr bins, except the poorly sampled Early Cretaceous 136–131 Myr bin. OPC values for dietary classes in extant rodents and carnivorans¹⁸ were mapped on Fig. 2a. Though there is some overlap in dietary classes, lower tooth row OPC values between 60 and 110 include mostly carnivores, between 110 and 160 mostly animal-dominated omnivores, between 160 and 200 mostly plant-dominated omnivores, and above 200 mostly herbivores.

2. Generic richness data collection

Patterns of generic richness provide a view of taxonomic diversification that can be decoupled from morphological diversification²³. Taxonomic and temporal range data

were compiled from the most recent compendia^{3,11,32} and primary literature available up to November 16, 2010. We standardized temporal ranges to the recent Geologic Time Scale of Gradstein³³. Taxonomic and temporal range data are in Supplementary Table 3. Richness was evaluated at the level of genera not species under the assumption that genera provide a more consistent record of long-term trends in taxonomic richness than more volatile patterns recorded at the species level. The number of described paulchoffatiid multituberculate genera from the Late Jurassic Guimarota fauna¹⁰⁹ is likely artificially inflated: some named taxa are based entirely on upper dentition and may belong to named taxa that are based entirely on lower dentition and vice versa³. To adjust for this, we removed six genera that were based entirely on upper dentitions. The data were partitioned into 5-Myr bins. Although partitioning the data into smaller bins would produce higher temporal resolution patterns, low sampling intensity for some intervals and poor age control for some taxa made that impractical. We placed the lower boundary of the first bin at 171 Myr ago so as to have the desired effect of placing a later bin boundary at the K-Pg boundary (ca. 66 Myr ago) rather than having it span this temporal boundary and associated mass extinction event. Whenever the lower temporal range of a taxon fell on an upper bound for a bin, it was counted in the younger bin. For example, the temporal range of *Acheronodon* is 66–60 Myr ago and was thus counted in the 66–61 Myr bin and the 61–56 Myr bin but not the 71–66 Myr bin. The same logic was applied to the upper temporal range of a taxon that fell on a lower bound for a bin. The intensity of mammalian fossil sampling varies through the geologic time interval of this study (171–31 Myr ago) as a result of research emphases and availability of fossil-bearing

exposures. The patterns of generic richness should thus be viewed as long-term trends rather than placing too much emphasis on individual fluctuations. We excluded a 5-Myr temporal bin from the analyses because of notably poor sampling; the 136–131 Myr bin included only one recorded multituberculate genus. In Figure 2a-c, we spanned this sampling gap with a dashed line connecting the data points on either side, the 141–136 Myr bin and the 131–126 Myr bin. Generic richness data are in Supplementary Table 4.

3. Body mass data collection

Body size provides a second measure of ecomorphological diversification and is correlated with a variety of life history traits in extant mammals³⁴. Estimates of multituberculate body mass were initially calculated based on a formula presented in Legendre³⁵ that was derived from a database of extant therian mammals that includes rodents, primates, artiodactyls, perissodactyls, lipotyphlans, chiropterans, carnivorans, and marsupials. The formula is as follows: $\text{LN}(\text{body mass}) = 1.827 * \text{LN}(\text{lower m1 area}) + 1.81$. Because multituberculates are phylogenetically outside of Theria (e.g., ref. 3), application of this formula to multituberculates requires justification. McDermott et al.³⁶ applied this formula, along with other formulae based on cranial and postcranial measurements, to multituberculates. They found that the generalized formula of Legendre³⁵ produced estimates convergent with estimates from other formulae, and thus deemed it reasonable to apply this formula to estimate multituberculate body mass. We estimated body mass for multituberculate species rather than genera because congeneric species are often discriminated from each other based on dental dimensions or other size

correlates; generic-level body mass estimates would overlook this variation. We calculated lower m1 area for 156 multituberculate species based on m1 length and width measurements that were compiled from the primary literature and our own measurements using a Leica MZ9.5 binocular dissecting microscope and custom measuring stage that has the capability of reading to the nearest 0.001 mm. When more than one measurement was available for a taxon, mean values were used to calculate lower m1 area. In a few cases measurements were not published and casts or specimens were not available, so estimates were made from published figures.

Body mass estimates based on lower m1 area appear reasonable for small-bodied multituberculates but are larger than expected for large-bodied multituberculates (e.g., *Taeniolabis taoensis* body mass estimate > 100 kg); this implies that scaling of lower m1 area to body mass differs in multituberculates and the reference group (Theria). Because the rodent skull and body plan is similar in appearance, rodents have often been suggested as a suitable analog for multituberculates. On this basis, we investigated predictive formulae that use modern rodents as a reference group³⁷⁻³⁹. From the supplementary data of ref. 38, we generated a bivariate plot of skull length (SL) vs. upper tooth row length (UTRL) of modern rodents (n = 35) onto which we plotted the same measurements available in 10 multituberculate taxa (Supplementary Fig. 3). For multituberculates, UTRL is the sum of the upper P4, M1, and M2 lengths. More broadly among mammals, SL is viewed as a more reliable predictor of body mass than molar dimensions⁴⁰ and has been used to estimate body mass in Mesozoic mammaliaforms⁴¹. Derivatives of SL, such as mandible length, have also been successfully used to estimate

body mass in non-therian Mesozoic mammals⁴². Unfortunately few multituberculate taxa are known from complete skulls. SL measurements for 10 multituberculate taxa, including the large-bodied *Taeniolabis taoensis*, are mostly based on specimens from the Campanian of Mongolia and Paleocene of North America. We concluded that UTRL and SL have a similar scaling relationship within rodents and multituberculates, because the multituberculate data plotted among the rodent data. Thus, rodents are an appropriate reference group for estimating body mass in multituberculates.

To further constrain the nature of discrepancy between estimates from the general therian formula³⁵ and rodent formulae^{37,39}, we plotted body mass estimates of multituberculates based on lower m1 area³⁵ against estimates from other anatomical elements (SL³⁹, UTRL³⁹, lower tooth row length³⁷) available for 10 to 13 multituberculate taxa. As estimates based on lower m1 area exceeded ca. 1 kg they became increasingly larger than estimates based on other anatomical elements. Because SL takes into account a greater proportion of the animal's total length than either UTRL or LTRL, we would expect body mass estimates based on SL to be more accurate than those based on UTRL or LTRL. Thus, whenever possible, we used SL to estimate body mass according to the formula³⁹: $\text{Log (body mass)} = 3.488 * (\text{Log SL}) - 3.332$. Unfortunately, SL is not as widely available as UTRL or LTRL. For large-bodied taxa, LTRL estimates were less than expected based on skull length. UTRL estimates were consistently greater than expected based on skull length (Supplementary Fig. 4d). Thus, for species without skull length data, we developed a regression formula to predict the SL body mass estimate from the lower m1 area body mass estimate, the most readily available estimate (see

Supplementary Fig. 5). This approach provides more conservative estimates of body mass than provided from UTRL, LTRL, and m1 area formulae. Estimates for the smallest multituberculates (under 100 g based on m1 area formula) are slightly greater when using the m1-SL regression formula and estimates for remaining multituberculates (over 100g based on the m1 area formula) are slightly less than estimates from UTRL or m1 area formulae, producing a more compressed body size distribution overall. Supplementary Table 5 includes lower m1 area, SL, predicted body mass, and sources for measurements. In Figure 2c, body mass estimates from Supplementary Table 5 were plotted as horizontal lines, lengths of which correspond to the temporal range of the species. We calculated geometric mean and standard deviation of body masses for each 5-Myr bin, except the Early Cretaceous 136–131 Myr bin. Nevertheless, because only ten of the 156 multituberculate body size estimates are based on skulls, which provide a more reliable basis to estimate body size than teeth alone, the body size trends should be considered approximate.

4. Correlation of estimated body mass and OPC

Because of the relationship between metabolic rate and body size in mammals⁴³⁻⁴⁷, small-bodied mammals often cannot subsist on a strictly herbivorous diet of leaves low in nutritional value⁴⁸⁻⁵⁰. Small-bodied mammals generally require a diet of high-energy content foods, such as fruits and insects. To test whether dental complexity as measured by OPC followed this pattern, we investigated the correlation of OPC and estimated body mass. Because OPC values in the sample are skewed toward higher

values, we log transformed the data. We found that $\ln(\text{body mass})$ and $\ln(\text{OPC})$ are strongly correlated in the sampled multituberculates (Pearson correlation coefficient = 0.672, $p < 0.01$, $n = 40$; Supplementary Fig. 6) and thus generally consistent with this claim; smaller-bodied multituberculates tend to have low OPC values corresponding to carnivorous or animal-dominated omnivorous diets, whereas larger-bodied multituberculates have high OPC values indicative of herbivorous diets with a greater proportion of stems and leaves. However, as in modern mammals, there are exceptions to this general pattern. The small-bodied multituberculate *Chulsanbaatar vulgaris* (13.3 g) has a high dental complexity (OPC 171), whereas the larger-bodied *Neoliotomus ultimus* (3.84 kg) has a low dental complexity (OPC 130).

5. Resampling routine and permutation simulation

To investigate whether the temporal patterns of OPC and body mass are due to differences in sampling, OPC and body mass values were permuted among taxa. OPC was randomized at the generic level, while body mass was randomized at the species level. Both measures were natural log transformed. Data were binned in 5-Myr durations where all taxa that occur within that bin were sampled. As described above, taxa that have their first appearance at the bottom of a bin were excluded. The K-Pg boundary (66 Myr ago) was used as the date from which bin values were calculated. In both cases, the age range of each measured taxon was preserved while the measurement information was randomly assigned to an age range. Because the desired alpha, or level of significance, was set at 0.05 and to reduce volatility⁵¹, this procedure was repeated 5000 times,

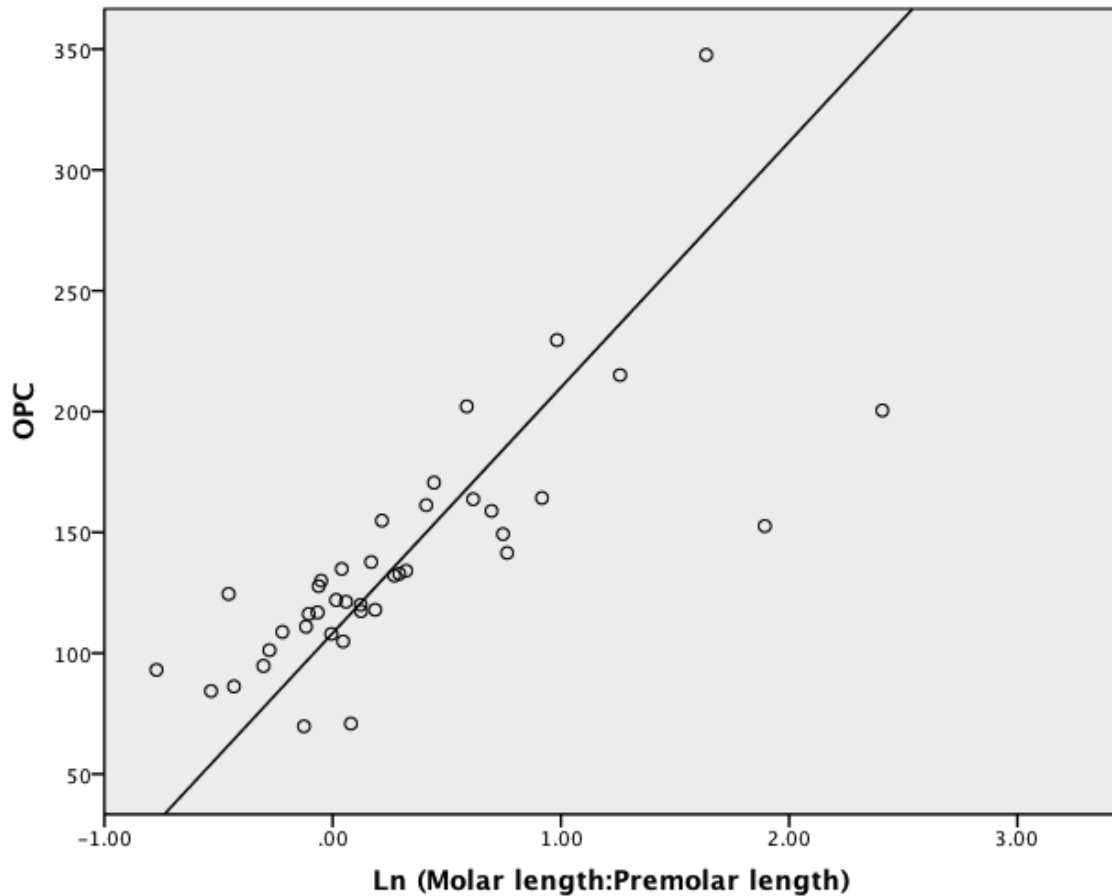
creating 5000 randomized mean values for each bin. All simulations were done in the R programming environment⁵². The random seed was set at the beginning of the simulation.

In Supplementary Figures 7 and 8 (OPC and body mass, respectively), the observed mean in every bin was plotted (black line) versus time and then compared to the 2.5 and 97.5 percentiles of the resampling distribution (dashed blue line). Significant difference is determined as when the observed mean value falls outside of this two-tailed confidence interval based on the resampled distribution (2.5–97.5 percentile).

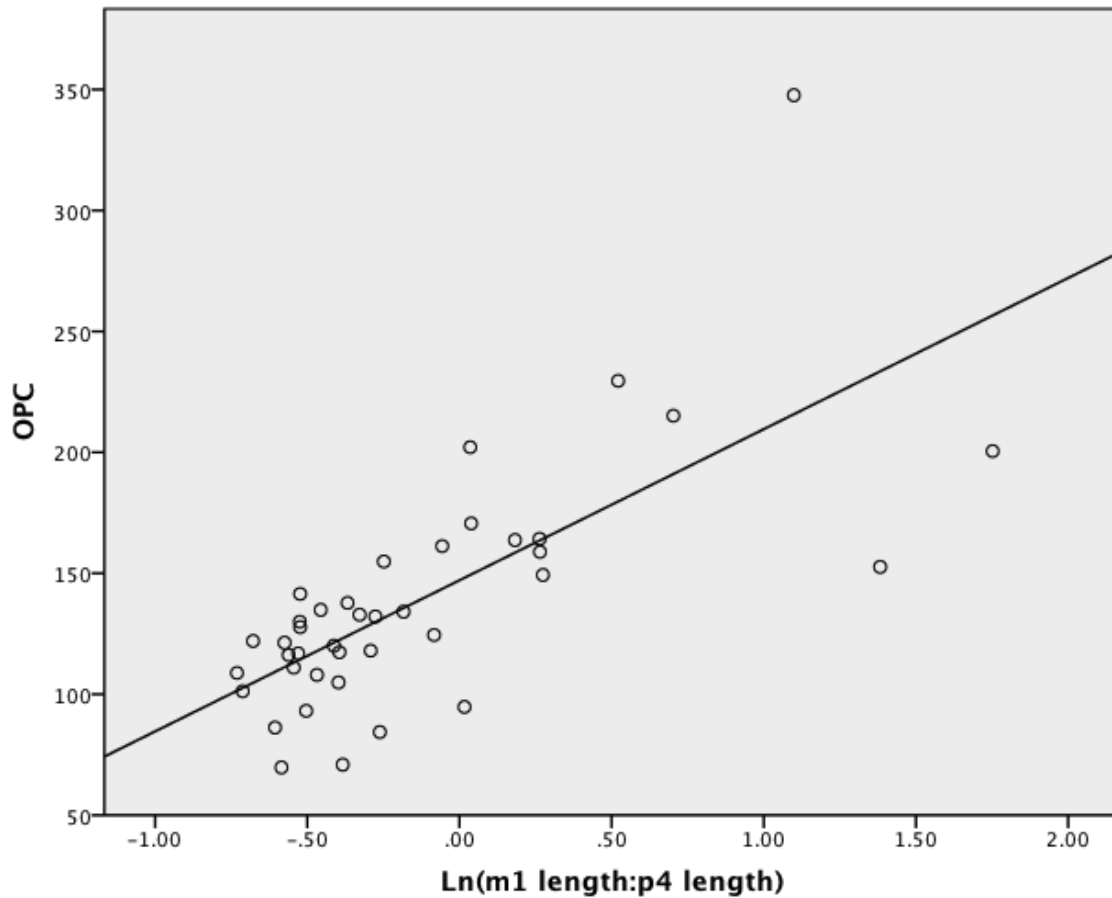
Observed OPC values fall outside of the two-tailed confidence interval in four bins (156–151, 146–141, 141–136 and 76–71 Myr ago). For 156–151, 146–141 and 141–136 Myr ago, the observed mean is significantly lower than the randomized distribution, and from 76–71 Myr ago the observed mean is significantly greater (Supplementary Fig. 7). The observed body mass values fall outside of the two-tailed confidence interval in four bins (146–141, 141–136, 101–96 and 71–66 Myr ago). In the first three of these bins, the observed mean is significantly lower than the randomized distribution, and in the 71–66 Myr bin the observed mean is significantly greater (Supplementary Fig. 8). In all other cases, the observed mean is not statistically different from the randomized distribution.

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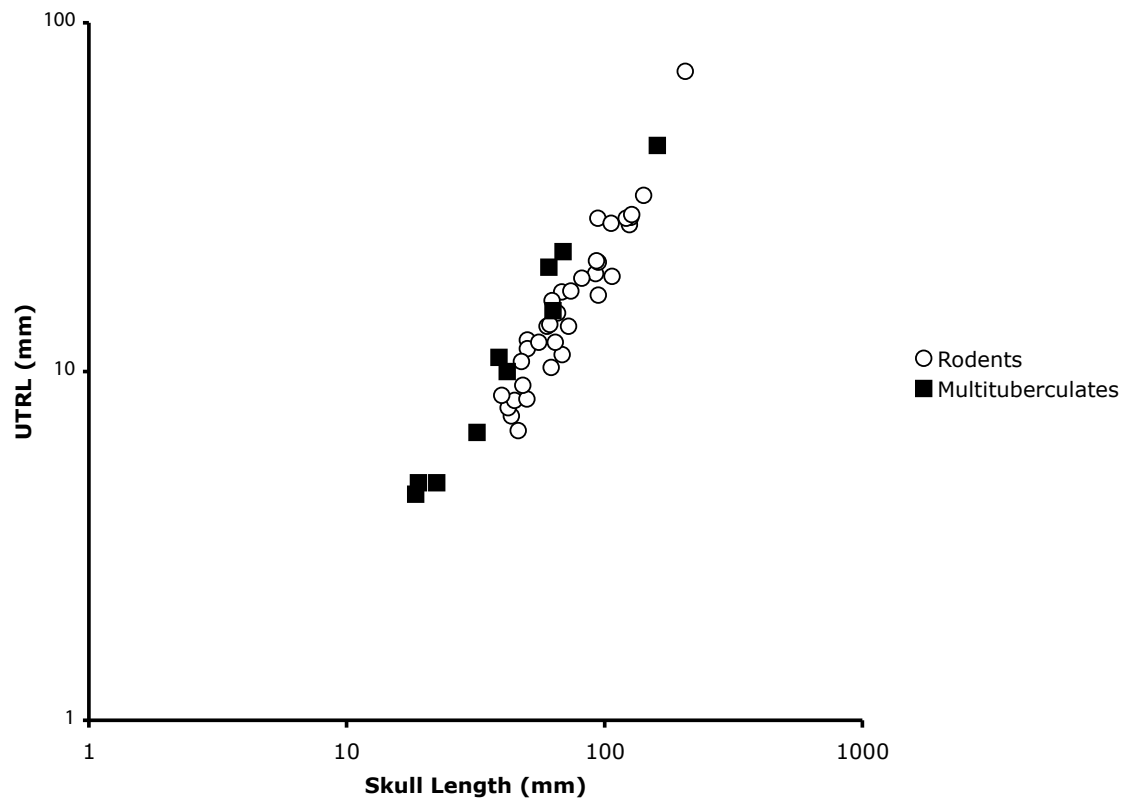
Supplementary Figure 1. Least-squares linear regression of OPC and the natural log of the lower molars to premolars length ratio. $R = 0.750$, $p < 0.0001$, $n = 40$. Predictive equation: $OPC = 121.44 + 57.20 * (\ln \text{ molars:premolars length ratio})$.



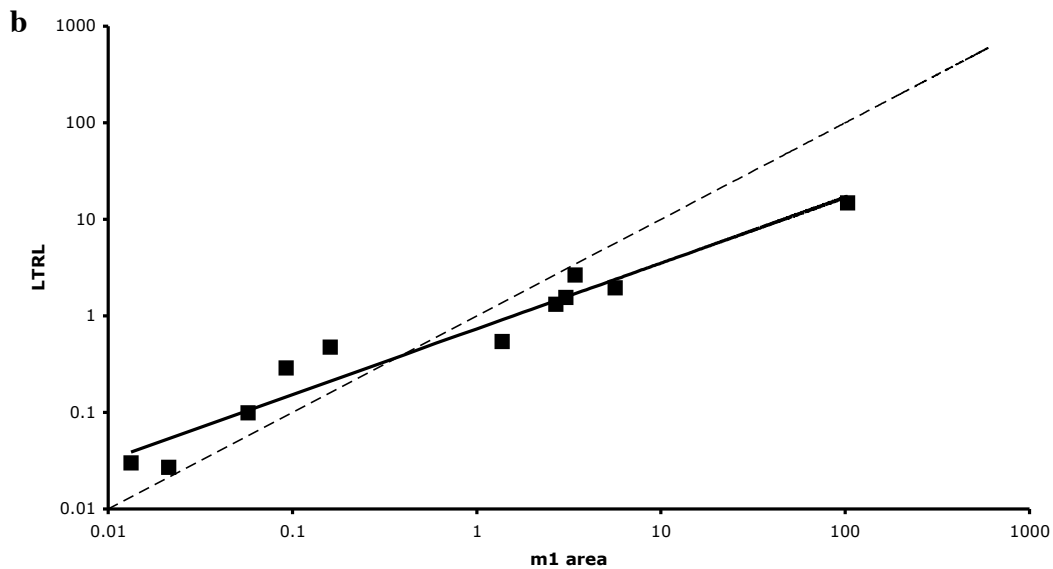
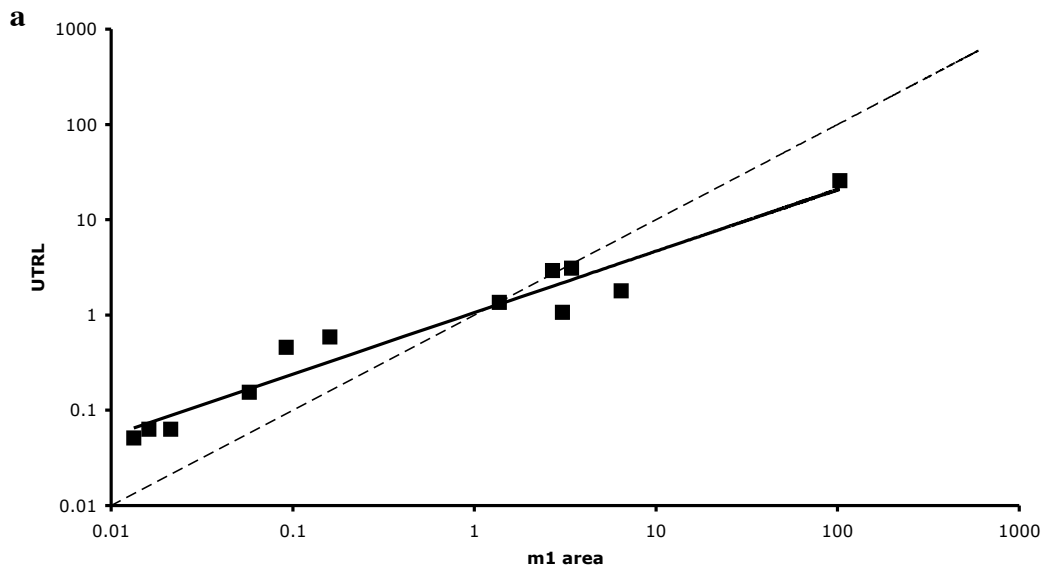
Supplementary Figure 2. Least-squares linear regression of OPC and the natural log of lower first molar (m1) to lower fourth premolar (p4) length ratio. $R = 0.711$, $p < 0.0001$, $n = 40$. Predictive equation: $OPC = 147.15 + 62.48 * (\ln \text{ m1:p4 length ratio})$.



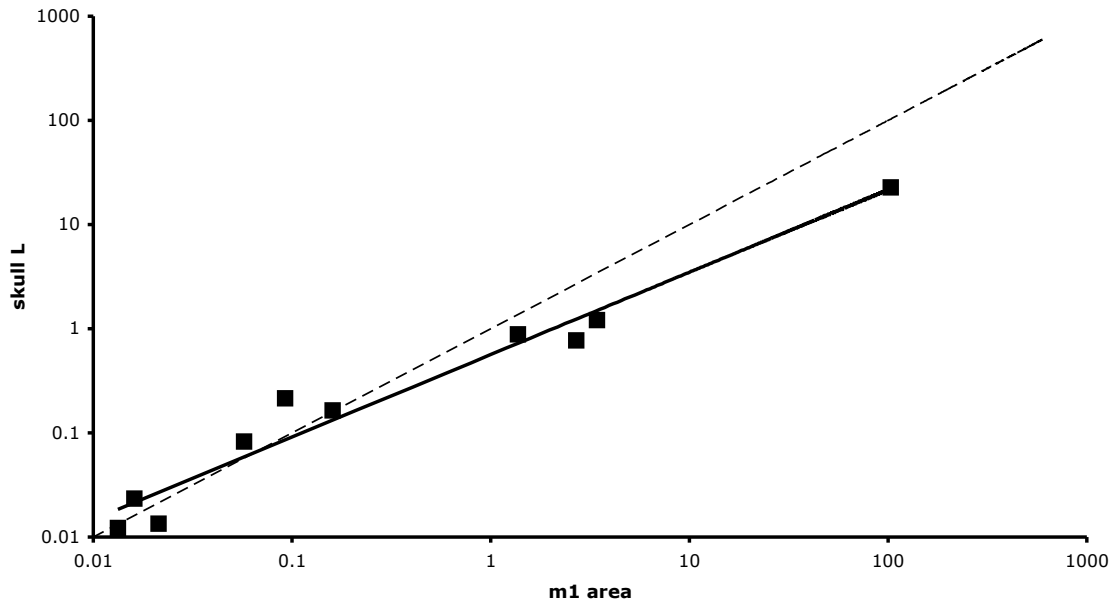
Supplementary Figure 3. Plot of skull length (SL) vs. upper tooth row length (UTRL) for rodent sample (open circles, $n = 35$) from ref. 38 and multituberculates (filled squares, $n = 35$) from this study.



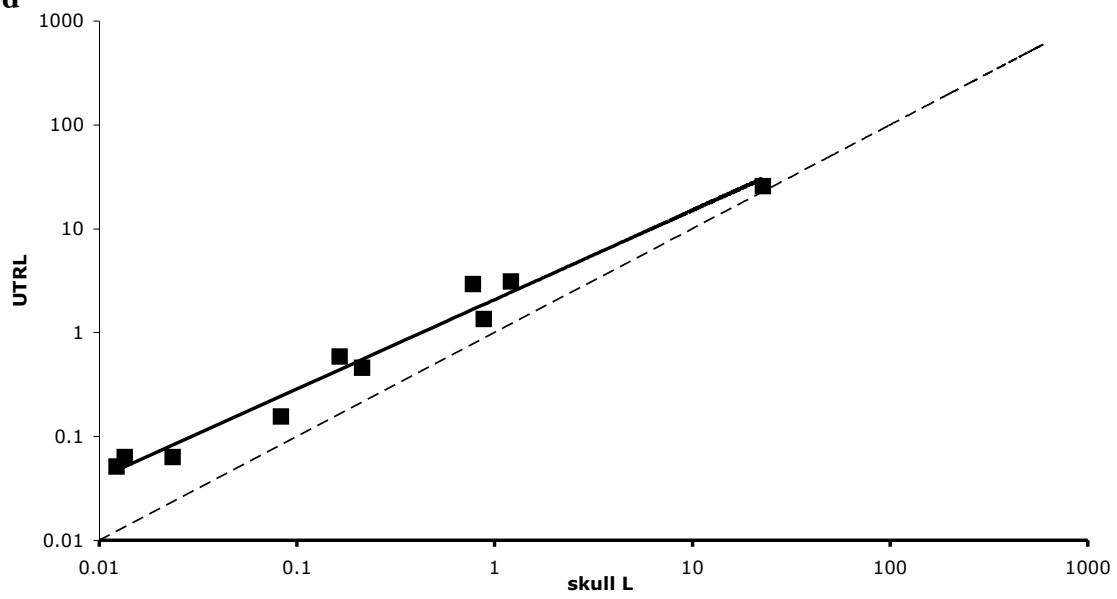
Supplementary Figure 4. Plots of body mass estimates based on lower first molar (m1) area³⁵ vs. (a) body mass estimates from upper tooth row length (UTRL³⁹, n = 12), (b) lower tooth row length (LTRL³⁷, n = 11), and (c) skull length (skull L³⁹, n = 10) for select multituberculate species; and (d) estimates from skull length vs. UTRL (n = 10). Scale in kilograms. Dashed line is $y=x$.



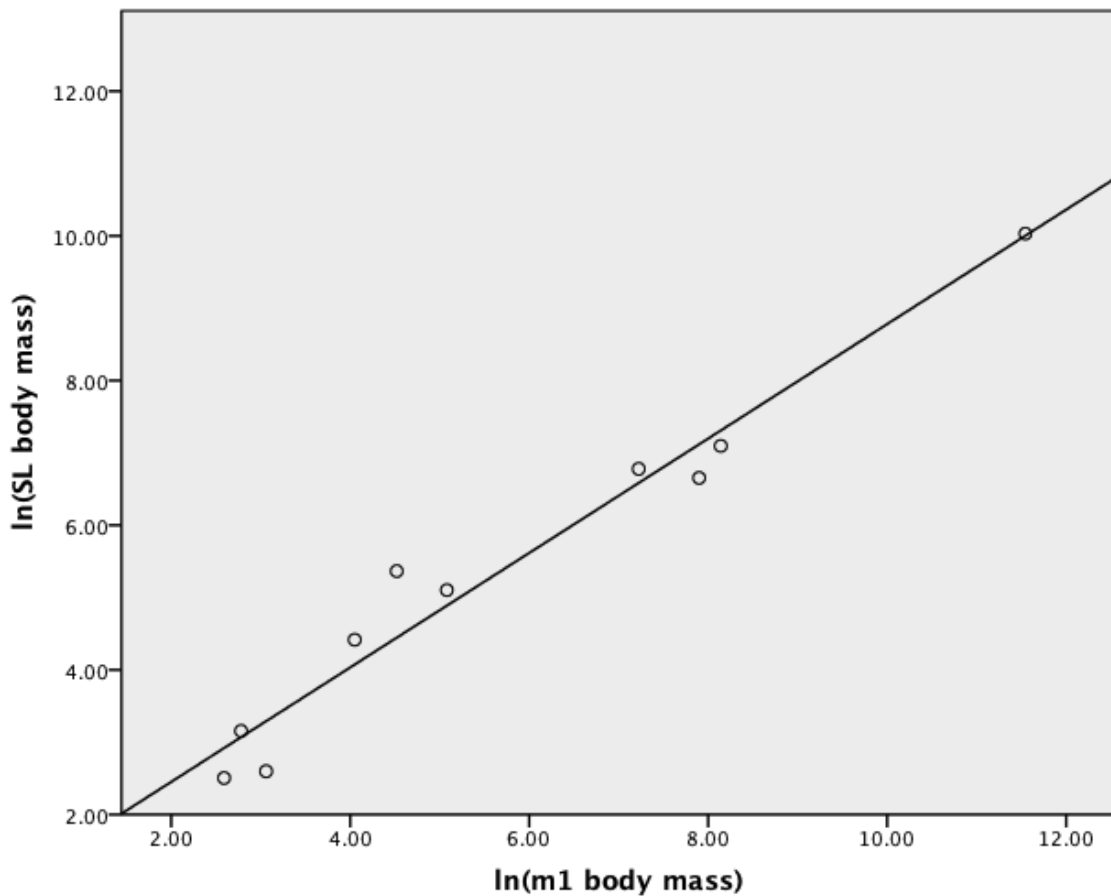
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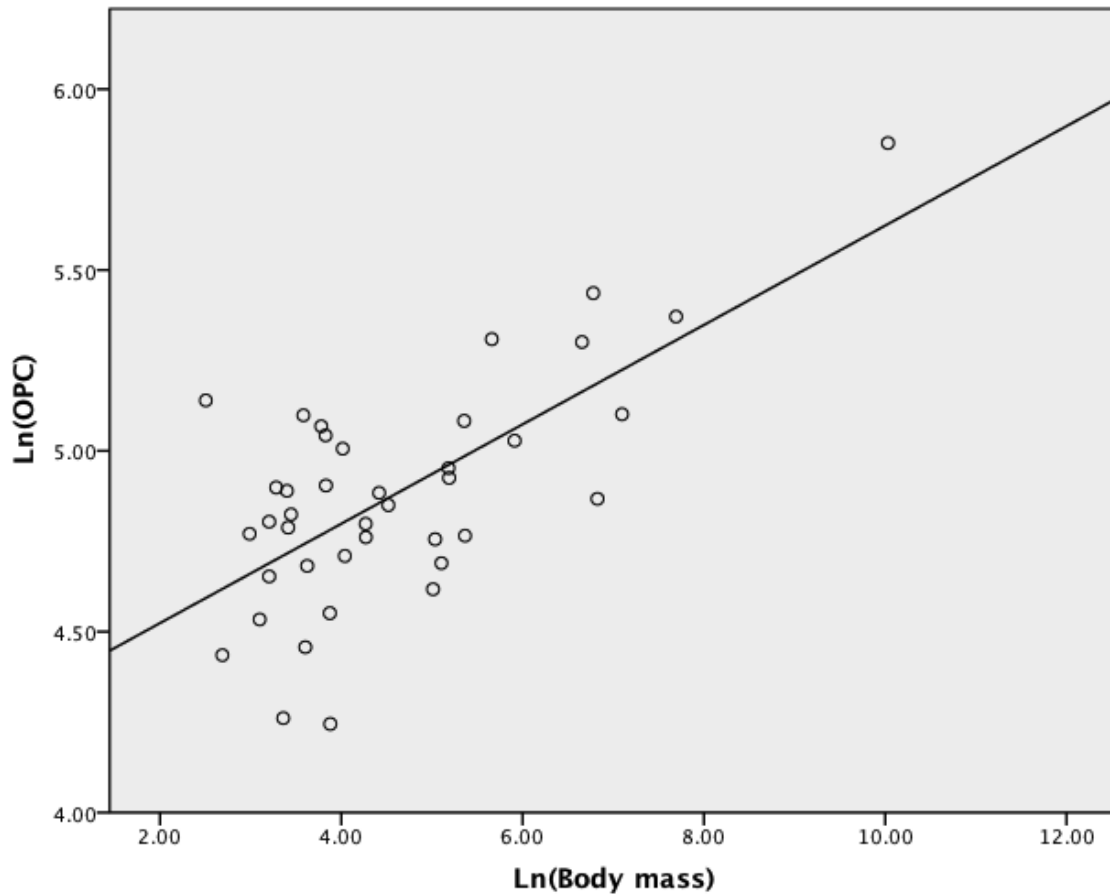
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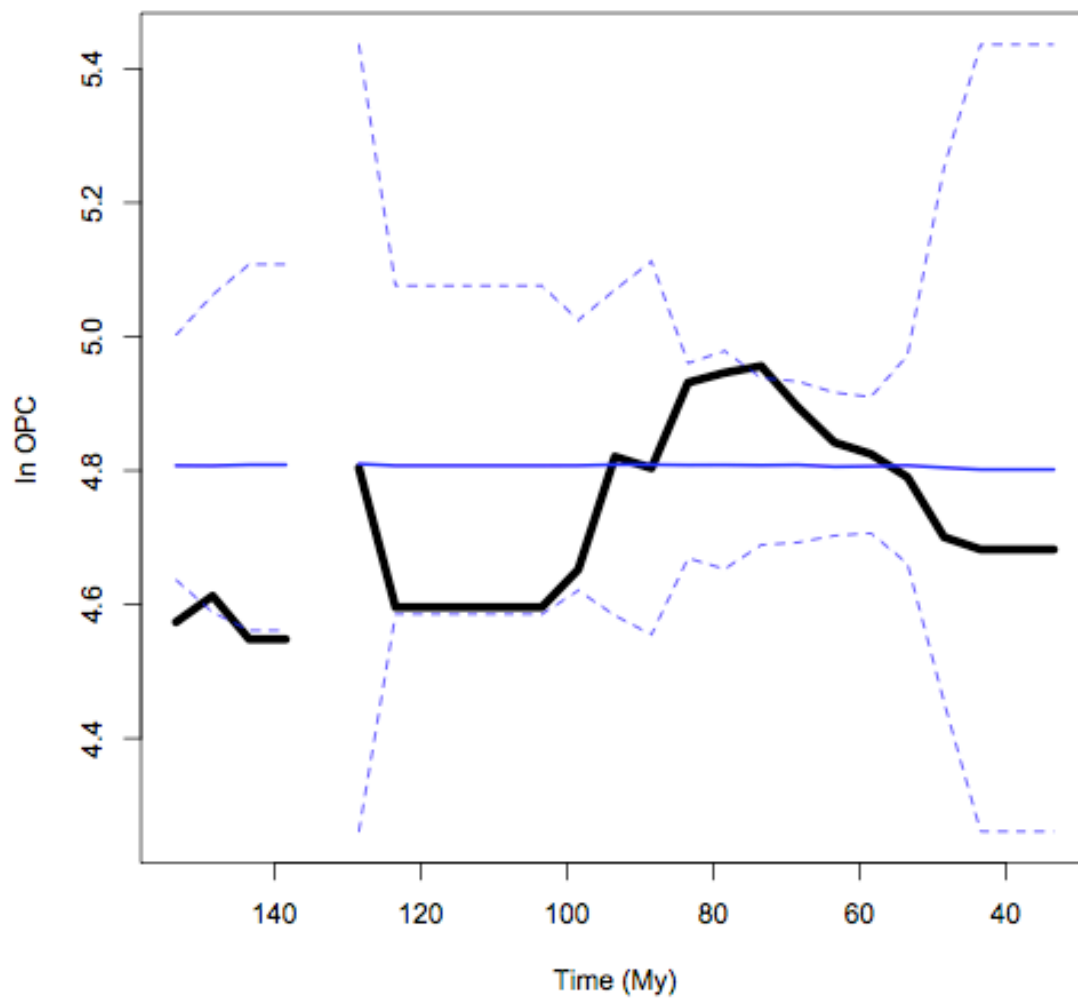
Supplementary Figure 5. Least-squares linear regression of the natural log of body mass estimated from lower first molar (m1) area³⁵ and the natural log of body mass estimated from the skull length (SL)³⁹ for select multituberculate species. $R = 0.980$, $p < 0.0001$, $n = 10$. Predictive equation: $\text{SL body mass estimate} = 0.87 + 0.79 * (\ln \text{ m1 body mass estimate})$.



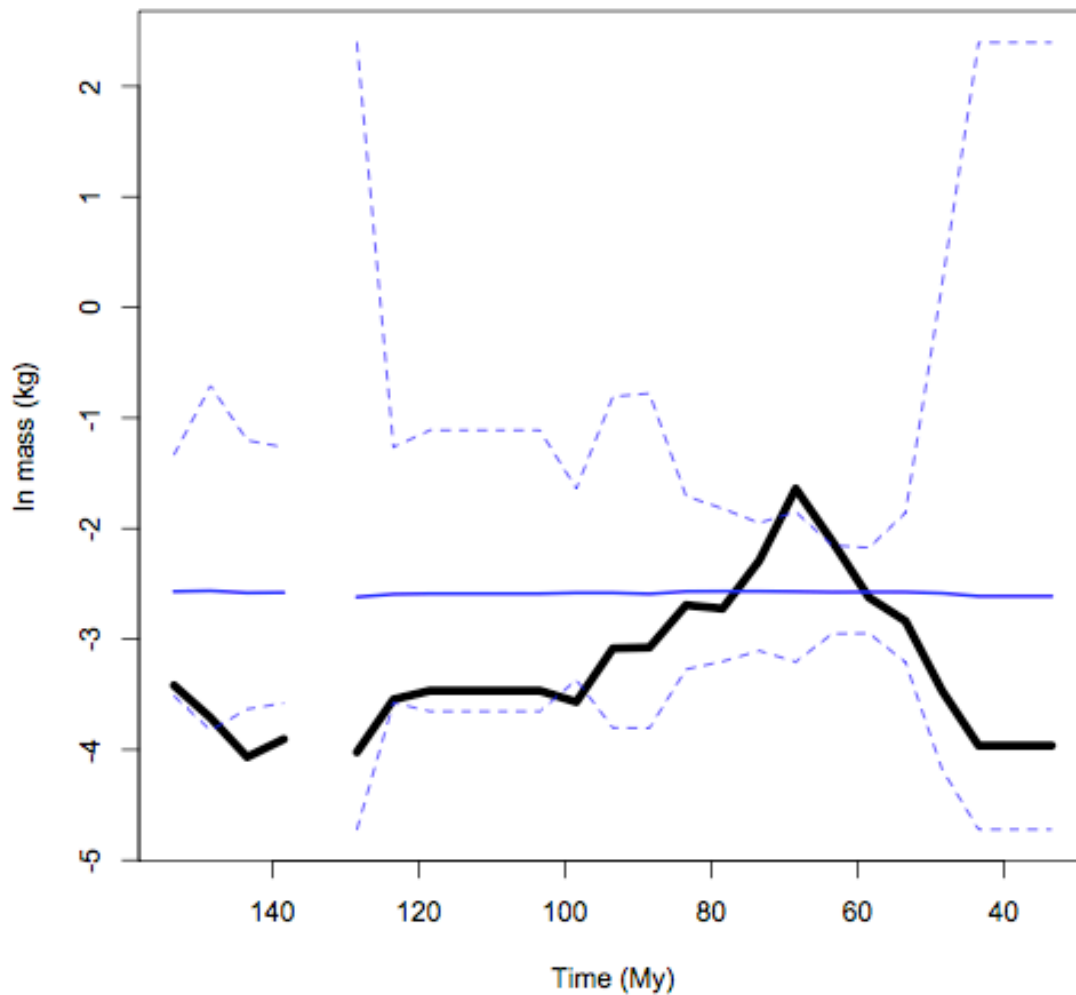
Supplementary Figure 6. Bivariate plot of the natural log of body mass estimates and natural log of OPC values for sampled multituberculates. Pearson correlation coefficient = 0.683, $p < 0.01$, $n = 40$.



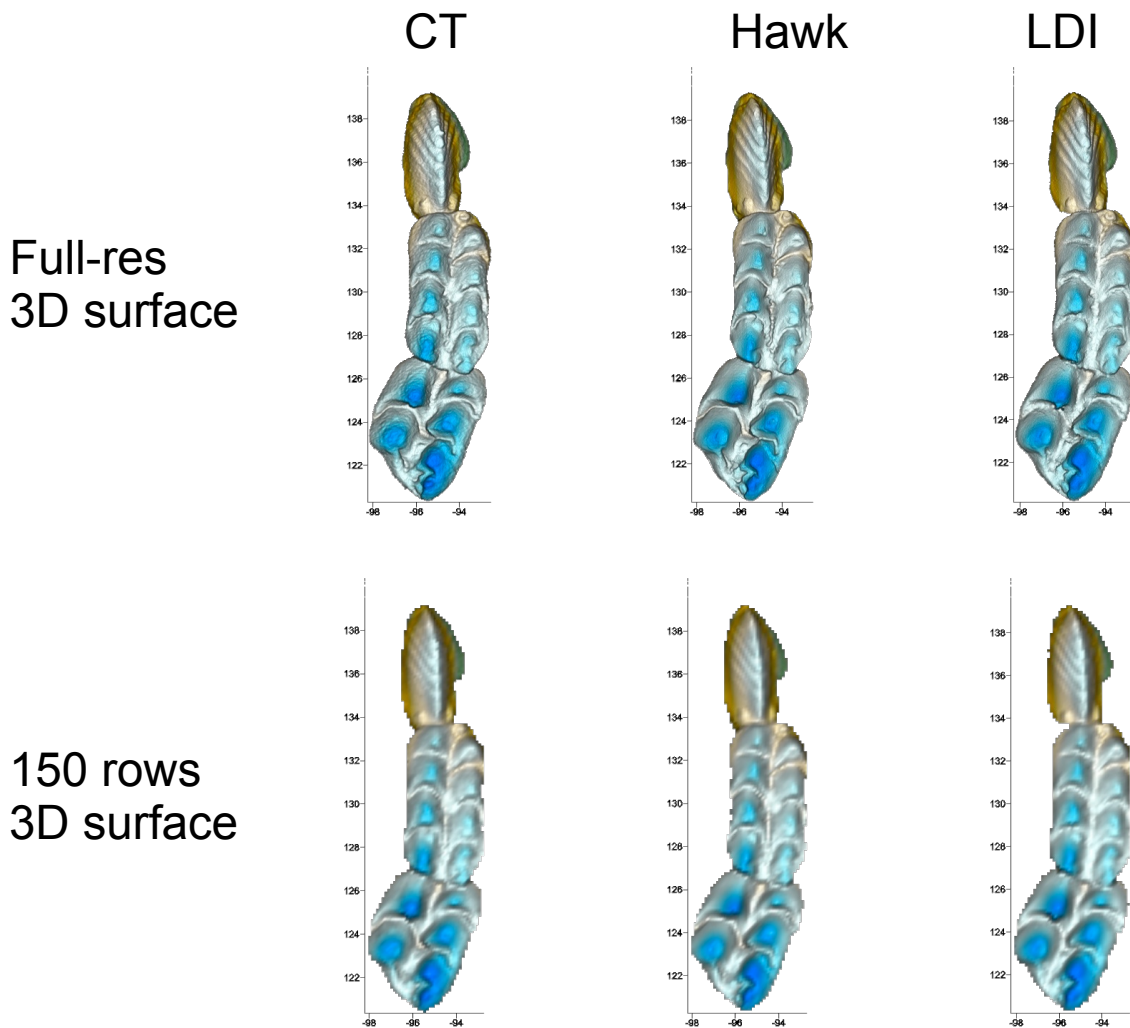
Supplementary Figure 7. Plot of natural log of OPC resampling simulation with mean (blue line) and 95% confidence limits (blue dashed lines) shown along with the mean of the natural log of the observed OPC (black line).



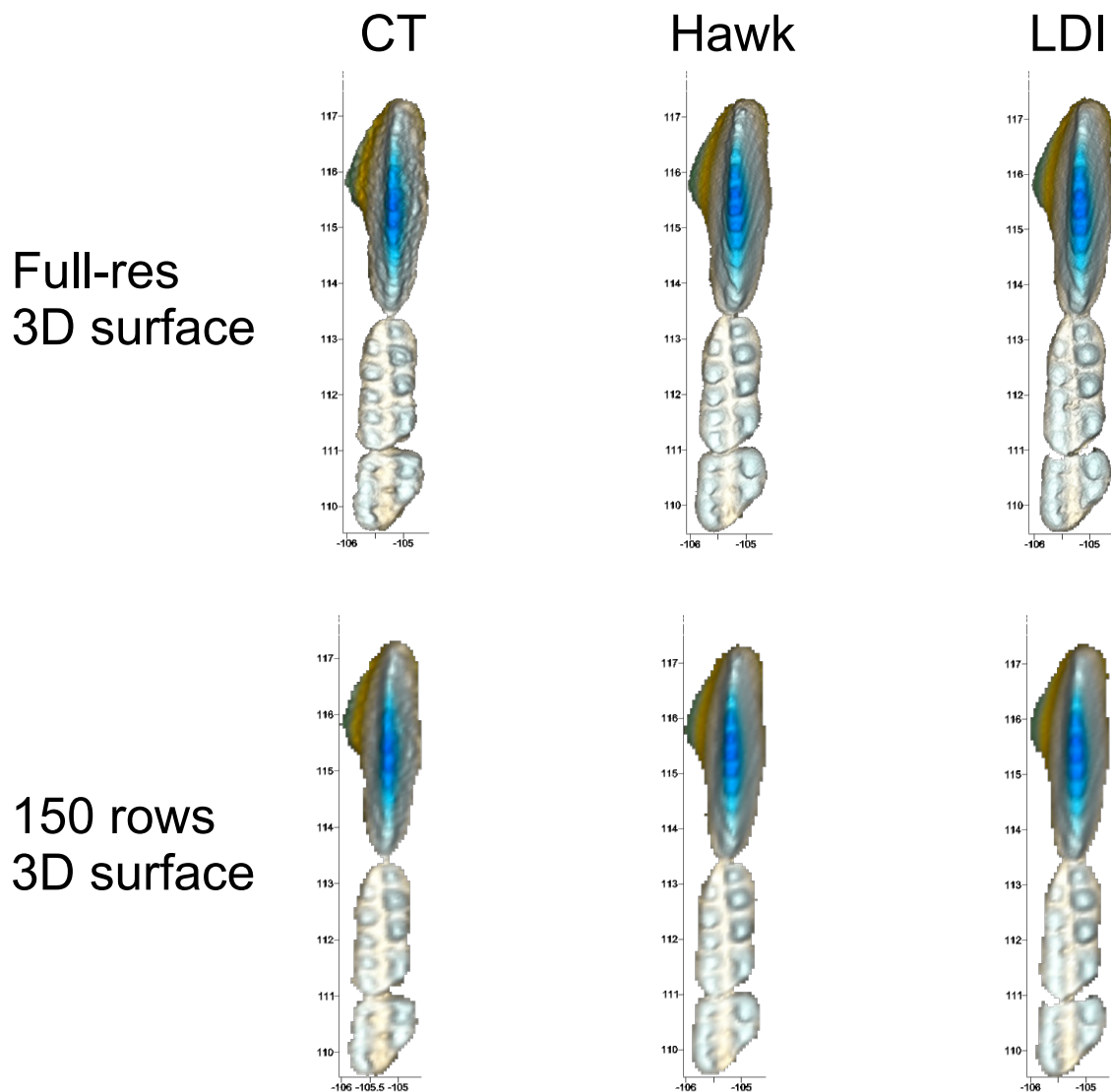
Supplementary Figure 8. Plot of natural log of body mass resampling simulation with mean (blue line) and 95% confidence limits (blue dashed lines) shown along with the mean of the natural log of observed body mass (black line).



Supplementary Figure 9. Comparisons of surfaces from three scanners scanning the same cast (*Meniscoessus robustus* UCMP 107405), illustrating both the full resolution surface and downsampled to 150 rows, as used in OPC analysis.



Supplementary Figure 10. Comparisons of surfaces from three scanners scanning the same cast (*Parikimys carpenteri* DMNH 52224), illustrating both the full resolution surface and downsampled to 150 rows, as used in OPC analysis.



Supplementary Table 1. Multituberculate specimens scanned, digitized, processed, and analyzed for OPC measurement. Each row represents a complete cheek tooth row, either from a single specimen or a composite reconstructed from multiple specimens, that was sampled in this study. OPC species is the average OPC value if the species has more than one sampled cheek tooth row. OPC genus is the average OPC value if the genus has more than one sampled cheek tooth row. Specimens in bold have moderate/heavy wear.

Species	Specimen #	OPC	OPC species	OPC genus
<i>Acheronodon vossae</i>	UALVP 24544, 42798, 24578	163.75	163.75	163.75
<i>Allocosmodon woodi</i>	UALVP 40494	154.875	154.875	154.875
<i>Anconodon cochranensis</i>	USNM 9765	111	111	111
<i>Baiotomeus douglassi</i>	USNM V9795	116.25	116.25	116.25
<i>Barbatodon transylvanicus</i> , <i>B. sp.</i> indet.	FGGUB R.1635, 1623	141.5	141.5	141.5
<i>Bolodon osborni</i>	BMNH 48399, DORCM GS2, 206	93.125	93.125	93.125
<i>Bryceomys fumosus</i>	MNA V7476, V6298, V6765	114	114	120.0625
<i>Bryceomys intermedius</i>	OMNH 34005, 33001, 26626	126.125	126.125	-
<i>Catopsalis alexanderi</i>	UCM 34979	215.125	215.125	215.125
<i>Catopsbaatar catopsaloides</i>	PM 120/107	202.625	202.625	202.625

Species	Specimen #	OPC	OPC species	OPC genus
<i>Cedaromys bestia</i>	OMNH 26636, 25752, 33186	122	122	121.3125
<i>Cedaromys parvus</i>	OMNH 32987, 25750, 25760	120.625	120.625	-
<i>Chulsanbaatar vulgaris</i>	ZPAL MgM-I/108	170.625	170.625	170.625
<i>Cimexomys judithae</i>	MOR 302	134.125	134.125	134.125
<i>Cimolodon nitidus</i>	AMNH 57860	137.75	137.75	137.75
<i>Cimolomys gracilis</i>	UCMP 51514, 51552, 51669	202.125	202.125	202.125
<i>Ctenacodon serratus</i>	YPM 11832	84.375	84.375	84.375
<i>Dakotamys malcolmi</i>	MNA V6056, V6384, V6488	132.875	132.875	132.875
<i>Ectypodus musculus</i>	AMNH 17391	100.625	100.625	108
<i>Ectypodus tardus</i>	PU 13265	115.375	115.375	-
<i>Eobaatar magnus</i>	PIN 3101/60, PIN 3101/53	122	122	122
<i>Glirodon grandis</i>	LACM 120452	124.5	124.5	124.5
<i>Hainina belgica</i> , <i>H. godfriauxi</i>	HIN 16-16 60-1, HIN 17-17 70-2, HIN 17-17 70-3	116.5	116.5	116.5
<i>Kimbetohia mziae</i>	UCM 38858	116.875	116.875	116.875
<i>Kryptobaatar dashzevegi</i>	GI-PST 8-2	132.125	132.125	132.125
<i>Lambdopsalis bulla</i>	IVPP V7152.17	200.5	200.5	200.5
<i>Meketibolodon robustus</i>	IPFUB Gui Mam 89/76	94.75	94.75	94.75

Species	Specimen #	OPC	OPC species	OPC genus
<i>Meniscoessus robustus</i>	UCMP 107405	164.25	164.25	164.25
<i>Mesodma archibaldi</i>	MNA V7531	114.75	114.75	118
<i>Mesodma pygmaea</i>	AMNH 35298	121.25	121.25	-
<i>Microcosmodon conus</i>	PU 22316	158.25	158.875	158.875
<i>Microcosmodon conus</i>	UALVP 42711	159.5	-	-
<i>Mimetodon silberlingi</i>	USNM 9798	70.875	70.875	70.875
<i>Nemegtbaatar gobiensis</i>	ZPAL MgM-I/81	117.375	117.375	117.375
<i>Neoliotomus ultimus</i>	UW 10433, 10428, 6577, 10430	130	130	130
<i>Neoplagiaulax hunteri</i>	UALVP 9955, 9787, 11977	69.25	69.25	69.25
<i>Parectypodus trovessartianus</i>	AMNH 3026	127.75	127.75	127.75
<i>Parikimys carpenteri</i>	DMNH 52224	134.875	134.875	134.875
<i>Pentacosmodon bowensis</i>	UALVP 42806, 24650, 42800	160	160	149.3125
<i>Pentacosmodon pronus</i>	MCZ 20066	138.625	138.625	-
<i>Plagiaulax becklesii</i>	BMNH 47731, 47733	86.25	86.25	86.25
<i>Prionessus lucifer</i>	AMNH 21731	152.625	152.625	152.625
<i>Prochetodon taxus</i>	UM 71311	101.25	101.25	101.25
<i>Ptilodus kummae</i>	UALVP 10912, 9001, 10253	106.75	106.75	108.8125
<i>Ptilodus montanus</i>	USNM 6076	110.875	110.875	-

Species	Specimen #	OPC	OPC species	OPC genus
<i>Stygmymys kuszmauli</i>	UCMP 92525, 92528, 133000	161.25	161.25	161.25
<i>Taeniolabis taoensis</i>	NMMNH P-8631	347.625	347.625	347.625
<i>Xyronomys robinsoni</i>	UCM 34975	104.875	104.875	104.875

Museum abbreviations: AMNH, American Museum of Natural History; BMNH, British Museum of Natural History; DMNH, Denver Museum of Nature & Science; FGGUB, Faculty of Geology and Geophysics, University of Bucharest, Romania; GI-PST and PM, Mongolian Academy of Sciences; HIN, Paleontology collections of l'Université des Sciences et Techniques de Montpellier, France; IPFUB, Instituts für Paläontologie der Freien Universität; IVPP, Institute of Vertebrate Paleontology and Paleoanthropology; LACM, Los Angeles County Museum; MCZ, Harvard University Museum of Comparative Zoology; MNA, Museum of Northern Arizona; MOR, Museum of the Rockies; NMMNH, New Mexico Museum of Natural History; OMNH, Oklahoma Museum of Natural History; PIN, Paleontological Institute of the Russian Academy of Sciences; PU, Princeton University collection held by Yale University Peabody Museum; UALVP, University of Alberta Vertebrate Paleontology collections; UCM, University of Colorado Museum; UCMP, University of California Museum of Paleontology; UM, University of Michigan Museum of Paleontology; USNM, United States National Museum; UW, University of Wyoming Geological Museum; YPM, Yale University Peabody Museum; ZPAL, Polish Academy of Sciences, Institute of Paleobiology.

Supplementary Table 2. Estimated OPC values for multituberculate species unavailable for this study. Predictive formulae are based on the regression analyses in Supplementary Figs. 1, 2.

Species	ln (molar L:premolar L)	ln (m1 L:p4 L)	estimated OPC
<i>Arginbaatar dimitrievae</i>	-0.775	-	77.13
<i>Buginbaatar transaltaiensis</i>	0.833	-	169.08
<i>Djadochtatherium matthewi</i>	0.608	-	156.23
<i>Essonodon browni</i>	0.221	-	134.11
<i>Eucosmodon americanus</i>	-	-0.638	107.28
<i>Guimarotodon leiriensis</i>	-0.626	-	85.62
<i>Heishanobaatar triangulus</i>	-0.317	-	103.30
<i>Iberodon quadrituberculatus</i>	0.057	-	124.71
<i>Kamptobaatar kuczynskii</i>	-0.288	-	104.98
<i>Krauseia clemensi</i>	0.037	-	123.54
<i>Kuehneodon dietrichi</i>	-0.310	-	103.70

Species	ln (molar L:premolar L)	ln (m1 L:p4 L)	estimated OPC
<i>Liaobaatar changi</i>	-0.499	-	92.88
<i>Liotomus marshi</i>	-0.461	-	95.09
<i>Mesodmops dawsonae</i>	-0.164	-	112.06
<i>Nessovbaatar multicostatus</i>	-0.121	-	114.49
<i>Nidimys occultus</i>	-	-0.300	128.41
<i>Paracimexomys priscus</i>	0.159	-	130.55
<i>Paulchoffatia delgadoi</i>	-0.510	-	92.25
<i>Pinheirodon pygmaeus</i>	-0.734	-	79.44
<i>Psalodon marshi</i>	-0.525	-	91.40
<i>Sinobaatar xiei</i>	-0.274	-	105.75
<i>Sloanbaatar mirabilis</i>	-	-0.463	118.24
<i>Uzbekbaatar wardi</i>	-	-0.591	110.21
<i>Zofiabaatar pulcher</i>	-	-0.685	104.32

Supplementary Table 3. Multituberculate genera with taxonomic, first appearance, last appearance, and geographic data.

Genera from the suborder Cimolodonta (boldface), genera from the suborder “Plagiaulacida” (not boldface). J = Jurassic, K = Cretaceous, As = Asia, Au = Australia, Eu = Europe, NA = North America, SA = South America, e = early, m = middle, l = late.

Genus	Superfamily	Family	FAD- Epoch	FAD- Age	FAD- Land Mammal Age	FAD- Myr	LAD- Epoch	LAD- Age	LAD- Land Mammal Age	LAD- Myr	Conti nent
<i>Hahnotherium</i>	-	Hahnotheriidae	J Mid	l. Bathon.	-	166	J Mid	l. Bathon.	-	165	Eu
<i>Kermackodon</i>	-	Kermackodontidae	J Mid	l. Bathon.	-	166	J Mid	l. Bathon.	-	165	Eu
<i>Bathmochoffatia</i>	Paulchoffatiid line	Paulchoffatiidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Guimarotodon</i>	Paulchoffatiid line	Paulchoffatiidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Henkelodon</i>	Paulchoffatiid line	Paulchoffatiidae	J Late	e.	-	155	J Late	l.	-	151	Eu

	d line	iidae		Kimmer.				Kimmer.			
<i>Kielanodon</i>	Paulchoffatii d line	Paulchoffat iidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Meketibolodon</i>	Paulchoffatii d line	Paulchoffat iidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Meketichoffatia</i>	Paulchoffatii d line	Paulchoffat iidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Paulchoffatia</i>	Paulchoffatii d line	Paulchoffat iidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Plesiochoffatia</i>	Paulchoffatii d line	Paulchoffat iidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Proalbionbaatar</i>	Plagiaulacid line	Albionbaat aridae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Pseudobolodon</i>	Paulchoffatii d line	Paulchoffat iidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Renatodon</i>	Paulchoffatii d line	Paulchoffat iidae	J Late	e. Kimmer.	-	155	J Late	l. Kimmer.	-	151	Eu
<i>Xenachoffatia</i>	Paulchoffatii	Paulchoffat	J Late	e.	-	155	J Late	l.	-	151	Eu

	d line	iidae		Kimmer.				Kimmer.			
<i>Ctenacodon</i>	Allodontid line	Allodontid ae	J Late	e. Kimmer.	-	155	J Late	e. Tithon.	-	147	NA
<i>Glirodon</i>	Allodontid line	-	J Late	e. Kimmer.	-	155	J Late	e. Tithon.	-	147	NA
<i>Morrisonodon</i>	Allodontid line	Allodontid ae	J Late	e. Kimmer.	-	155	J Late	e. Tithon.	-	147	NA
<i>Psalodon</i>	Allodontid line	Allodontid ae	J Late	e. Kimmer.	-	155	J Late	e. Tithon.	-	147	NA
<i>Zofiabaatar</i>	Allodontid line	Zofiabaatar idae	J Late	e. Kimmer.	-	155	J Late	e. Tithon.	-	147	NA
<i>Kuehneodon</i>	Paulchoffatii d line	Paulchoffat iidae	J Late	e. Kimmer.	-	155	J Late	l. Tithon.	-	146	Eu
<i>Albionbaatar</i>	Plagiaulacid line	Albionbaat aridae	K Early	e. Berrias.	-	146	K Early	l. Berrias.	-	140	Eu
<i>Bernardodon</i>	Paulchoffatii d line	Pinheirodo ntidae	K Early	e. Berrias.	-	146	K Early	l. Berrias.	-	140	Eu
<i>Bolodon</i>	Plagiaulacid	Plagiaulaci	K	e.	-	146	K Early	l. Berrias.	-	140	Eu

	line	dae	Early	Berrias.							
<i>Ecprepaulax</i>	Paulchoffatii d line	Pinheirodo ntidae	K Early	e. Berrias.	-	146	K Early	l. Berrias.	-	140	Eu
<i>Gerhardodon</i>	Paulchoffatii d line	Pinheirodo ntidae	K Early	e. Berrias.	-	146	K Early	l. Berrias.	-	140	Eu
<i>Iberodon</i>	Paulchoffatii d line	Pinheirodo ntidae	K Early	e. Berrias.	-	146	K Early	l. Berrias.	-	140	Eu
<i>Pinheirodon</i>	Paulchoffatii d line	Pinheirodo ntidae	K Early	e. Berrias.	-	146	K Early	l. Berrias.	-	140	Eu
<i>Plagiaulax</i>	Plagiaulacid line	Plagiaulaci dae	K Early	e. Berrias.	-	146	K Early	l. Berrias.	-	140	Eu
<i>Sunnyodon</i>	Paulchoffatii d line	Paulchoffat iidae	K Early	e. Berrias.	-	146	K Early	l. Berrias.	-	140	Eu
<i>Loxaulax</i>	Plagiaulacid line	Eobaatarid ae	K Early	e. Valangin	-	140	K Early	e. Valangin.	-	139	Eu
<i>Cantalera</i>	Paulchoffatii d line	Pinheirodo ntidae	K Early	e. l. Hauter.	-	134	K Early	e. Barrem.	-	128	Eu

<i>Lavocatia</i>	Paulchoffatii d line	Pinheirodo ntidae	K Early	e. Barrem.	-	130	K Early	e. Barrem.	-	128	Eu
<i>Parendotheriu m</i>	Plagiaulacid line	Eobaatarid ae	K Early	e. Barrem.	-	130	K Early	e. Barrem.	-	128	Eu
<i>Galveodon</i>	Paulchoffatii d line	Paulchoffat iidae	K Early	e. Barrem.	-	130	K Early	l. Barrem.	-	125	Eu
<i>Hakusanobaat ar</i>	Plagiaulacid line	Eobaatarid ae	K Early	e. Barrem.	-	130	K mid	e. Aptian	-	121	As
<i>Tedoribaatar</i>	Plagiaulacid line	Eobaatarid ae	K Early	e. Barrem.	-	130	K mid	e. Aptian	-	121	As
<i>Eobaatar</i>	Plagiaulacid line	Eobaatarid ae	K Early	e. Barrem.	-	130	K mid	l. Albian	-	100	As
<i>Corriebaatar</i>	-	Corriebaat aridae	K mid	e. Aptian	-	125	K mid	l. Aptian	-	112	Au
<i>Arginbaatar</i>	-	Arginbaata ridae	K mid	e. Aptian	-	125	K mid	l. Albian	-	100	As
<i>Heishanobaat ar</i>	Plagiaulacid line	Eobaatarid ae	K Early	e. Aptian	-	125	K mid	l. Albian	-	100	As

<i>Kielanobaatar</i>	Plagiaulacid line	Albionbaataridae	K Early	e. Aptian	-	125	K mid	l. Albian	-	100	As
<i>Liaobaatar</i>	Plagiaulacid line	Eobaataridae	K mid	e. Aptian	-	125	K mid	l. Albian	-	100	As
<i>Monobaatar</i>	Plagiaulacid line	Eobaataridae	K mid	e. Aptian	-	125	K mid	l. Albian	-	100	As
<i>Sinobaatar</i>	Plagiaulacid line	Eobaataridae	K mid	e. Aptian	-	125	K mid	l. Albian	-	100	As
<i>Ameribaatar</i>	Paracimexomyids group	-	K mid	e. Cenoman	-	100	K mid	e. Cenoman	-	96	NA
<i>Janumys</i>	Plagiaulacid line	-	K mid	e. Cenoman	-	100	K mid	e. Cenoman	-	96	NA
<i>Bryceomys</i>	Paracimexomyids group	-	K mid	e. Cenoman	-	100	K mid	l. Turonian	-	89	NA
<i>Cedaromys</i>	Paracimexomyids group	-	K mid	e.	-	100	K mid	e.	-	74	NA

	ys group			Cenoman				Cenoman			
<i>Dakotamys</i>	Paracimexomys group	-	K mid	l. Cenoman	-	96	K mid	l. Cenoman	-	94	NA
<i>Cimolodon</i>	Ptilodontoidea	Cimolodontidae	K mid	l. Cenoman	-	96	K Late	l. Maastrich	Lancian	66	NA
<i>Uzbekbaatar</i>	-	-	K mid	e. Turon.	-	94	K mid	l. Coniac.	-	86	As
<i>Meniscoessus</i>	-	Cimolomyidae	K Late	e. Santon.	-	86	K Late	l. Maastrich	Lancian	66	NA
<i>Bulganbaatar</i>	Djadochtatheroidea	-	K Late	e. Campan.	-	84	K Late	e. Campan.	-	81	As
<i>Djadochtatherium</i>	Djadochtatheroidea	Djadochtatheriidae	K Late	e. Campan.	-	84	K Late	e. Campan.	-	81	As
<i>Kamptobaatar</i>	Djadochtatheroidea	Sloanbaataridae	K Late	e. Campan.	-	84	K Late	e. Campan.	-	81	As

<i>Sloanbaatar</i>	Djadochtathe roidea	Sloanbaata ridae	K Late	e. Campan.	-	84	K Late	e. Campan.	-	81	As
<i>Tombaatar</i>	Djadochtathe roidea	Djadochtat heriidae	K Late	e. Campan.	-	84	K Late	e. Campan.	-	81	As
<i>Viridomys</i>	-	-	K Late	e. Campan.	Aquilan	84	K Late	e. Campan.	Aquilan	81	NA
<i>Chulsanbaatar</i>	Djadochtathe roidea	-	K Late	e. Campan.	-	84	K Late	l. Campan.	-	71	As
<i>Kryptobaatar</i>	Djadochtathe roidea	Djadochtat heriidae	K Late	e. Campan.	-	84	K Late	l. Campan.	-	71	As
<i>Argentodites</i>	-	-	K Late	e. Campan.	-	84	K Late	l. Maastrich	-	66	SA
<i>Cimolomys</i>	-	Cimolomyi dae	K Late	e. Campan.	Aquilan	84	K Late	l. Maastrich	Lancian	66	NA
<i>Paracimexomys</i>	Paracimexom ys group	-	K mid	l. Santon.	-	84	K Late	l. Maastrich	Lancian	66	NA

<i>Cimexomys</i>	Paracimexomys group	-	K Late	e. Campan.	Aquilan	84	Paleogene	e. Paleoc.	l. Puercan	64	NA
<i>Mesodma</i>	Ptilodontoidea	Neoplagiatalacidae	K Late	e. Campan.	Aquilan	84	Paleogene	l. Paleoc.	m. Tiffania	57	NA
<i>Kaiparomys</i>	Ptilodontoidea	Cimolodontidae	K Late	m. Campan.	Judithian	79	K Late	l. Campan.	Judithian	74	NA
<i>Kimbetohia</i>	Ptilodontoidea	Ptilodontidae	K Late	m. Campan.	Judithian	79	Paleogene	e. Paleoc.	l. Puercan	64	NA
<i>Catopsbaatar</i>	Djadochtatheroidea	Djadochtatheriidae	K Late	l. Campan.	-	76	K Late	l. Campan.	-	71	As
<i>Nemegtbaatar</i>	Djadochtatheroidea	-	K Late	l. Campan.	-	76	K Late	l. Campan.	-	71	As
<i>Nessovbaatar</i>	Djadochtatheroidea	Sloanbaataridae	K Late	l. Campan.	-	76	K Late	l. Campan.	-	71	As
<i>Stygimys</i>	-	Eucosmodontidae	K Late	l. Campan.	-	75	Paleogene	e. Paleoc.	l. Torrejo	61	NA

									nian		
<i>Nidimys</i>	Ptilodontoidea	Neoplagiaticidae	K Late	l. Campan.	"Edmontonian"	74	K Late	e. Maastricht	"Edmontonian"	69	NA
<i>Barbatodon</i>	-	Kogaionidae	K Late	e. Maastricht	-	71	K Late	l. Maastricht	-	66	Eu
<i>Buginbaatar</i>	-	Cimolomyidae	K Late	e. Maastricht	-	71	K Late	l. Maastricht	-	66	As
<i>Bubodens</i>	Taeniolabidoidea	Taeniolabidae	K Late	l. Maastricht	Lancian	69	K Late	l. Maastricht	Lancian	66	NA
<i>Clemensodon</i>	-	Eucosmodontidae	K Late	l. Maastricht	Lancian	69	K Late	l. Maastricht	Lancian	66	NA
<i>Essonodon</i>	-	Cimolomyidae	K Late	l. Maastricht	Lancian	69	K Late	l. Maastricht	Lancian	66	NA

				h.				.			
<i>Kogaionon</i>	-	Kogaionidae	K Late	l. Maastric h.	-	69	K Late	l. Maastrich .	-	66	Eu
<i>Paressonodon</i>	-	Cimolomyidae	K Late	l. Maastric h.	Lancian	69	K Late	l. Maastrich .	Lancian	66	NA
<i>Parikimys</i>	Ptilodontoidea	Neoplagiulacidae	K Late	l. Maastric h.	Lancian	69	K Late	l. Maastrich .	Lancian	66	NA
<i>Hainina</i>	-	Kogaionidae	K Late	l. Maastric h.	-	69	Paleogene	l. Paleoc.	Thanetian	55	Eu
<i>Neoplagiulax</i>	Ptilodontoidea	Neoplagiulacidae	K Late	l. Maastric h.	Lancian	69	Paleogene	l. Paleoc.	Thanetian	55	NA
<i>Parectypodus</i>	Ptilodontoidea	Neoplagiulacidae	K Late	l. Maastric	Lancian	69	Paleogene	e. Eocene	l. Wasatchian	52	NA

				h.					hian		
<i>Xyromys</i>	Ptilodontoidea	Neoplagiulacidae	Paleogene	e. Paleoc.	e. Puercan	66	Paleogene	e. Paleoc.	m. Torrejonian	61	NA
<i>Acheronodon</i>	-	Microcosmodontidae	Paleogene	e. Paleoc.	e. Puercan	66	Paleogene	l. Paleoc.	e. Tiffanian	60	NA
<i>Catopsalis</i>	Taeniolabidoidea	Taeniolabididae	Paleogene	e. Paleoc.	e. Puercan	66	Paleogene	l. Paleoc.	m. Tiffanian	57	NA
<i>Ptilodus</i>	Ptilodontoidea	Ptilodontidae	Paleogene	e. Paleoc.	e. Puercan	66	Paleogene	l. Paleoc.	l. Tiffanian	56	NA
<i>Ectypodus</i>	Ptilodontoidea	Neoplagiulacidae	Paleogene	e. Paleoc.	e. Puercan	66	Paleogene	l. Eocene	m. Chadronian	35	NA
<i>Taeniolabis</i>	Taeniolabidoidea	Taeniolabididae	Paleogene	e. Paleoc.	m. Puercan	65	Paleogene	e. Paleoc.	l. Puercan	64	NA

<i>Eucosmodon</i>	-	Eucosmodontidae	Paleogene	e. Paleoc.	m. Puercan	65	Paleogene	e. Paleoc.	l. Torrejonian	61	NA
<i>Cernaysia</i>	Ptilodontoidea	Neoplagiulacidae	Paleogene	l. Paleoc.	-	65	Paleogene	l. Paleoc.	-	56	Eu, NA
<i>Microcosmodon</i>	-	Microcosmodontidae	Paleogene	e. Paleoc.	m. Puercan	65	Paleogene	l. Paleoc.	l. Clarkforkian	55	NA
<i>Xanclomys</i>	Ptilodontoidea	Neoplagiulacidae	Paleogene	e. Paleoc.	m. Torrejonian	62	Paleogene	e. Paleoc.	m. Torrejonian	61	NA
<i>Krauseia</i>	Ptilodontoidea	Neoplagiulacidae	Paleogene	e. Paleoc.	m. Torrejonian	62	Paleogene	l. Paleoc.	e. Tiffanian	60	NA
<i>Anconodon</i>	Ptilodontoidea	Cimolodontidae	Paleogene	e. Paleoc.	m. Torrejonian	62	Paleogene	l. Paleoc.	e. Tiffanian	59	NA
<i>Boffius</i>	-	Boffiidae	Paleogene	m.	-	62	Paleogene	m.	-	59	Eu

			ene	Paleoc.				Paleoc.			
<i>Baiotomeus</i>	Ptilodontoidea	Ptilodontidae	Paleogene	e. Paleoc.	l. Torrejonia	62	Paleogene	l. Paleoc.	m. Tiffania	58	NA
<i>Mimetodon</i>	Ptilodontoidea	Neoplagiailacidae	Paleogene	e. Paleoc.	l. Torrejonia	62	Paleogene	l. Paleoc.	l. Tiffania	56	NA
<i>Fractinus</i>	-	-	Paleogene	l. Paleoc.	e. Tiffanian	61	Paleogene	l. Paleoc.	e. Tiffania	60	NA
<i>Allocosmodon</i>	-	Microcosmodontidae	Paleogene	l. Paleoc.	e. Tiffanian	61	Paleogene	l. Paleoc.	m. Tiffania	57	NA
<i>Pentacosmodon</i>	-	Microcosmodontidae	Paleogene	l. Paleoc.	e. Tiffanian	61	Paleogene	l. Paleoc.	l. Tiffania	56	NA
<i>Liotomus</i>	Ptilodontoidea	Cimolodontidae	Paleogene	l. Paleoc.	e. Tiffanian	61	Paleogene	l. Paleoc.	Thanetian	55	NA/Eu

<i>Lambdopsalis</i>	Taeniolabidoi dea	Taeniolabi didae	Paleog ene	l. Paleoc.	-	59	Paleogene	l. Paleoc.	-	55	As
<i>Prionessus</i>	Taeniolabidoi dea	Taeniolabi didae	Paleog ene	l. Paleoc.	-	59	Paleogene	l. Paleoc.	-	55	As
<i>Prochetodon</i>	Ptilodontoide a	Ptilodontid ae	Paleog ene	l. Paleoc.	l. Tiffanian	59	Paleogene	l. Paleoc.	l. Clarkfo rkian	55	NA
<i>Sphenopsalis</i>	Taeniolabidoi dea	Taeniolabi didae	Paleog ene	l. Paleoc.	-	59	Paleogene	l. Paleoc.	-	55	As
<i>Neoliotomus</i>	Ptilodontoide a	-	Paleog ene	l. Paleoc.	l. Tiffanian	57	Paleogene	e. Eocene	m. Wasatc hian	53	NA
<i>Mesodmops</i>	Ptilodontoide a	Neoplagiau lacidiae	Paleog ene	l. Paleoc.	-	56	Paleogene	e. Eocene	-	50	As

Supplementary Table 4. Generic richness data from Supplementary Table 3 partitioned into 5-Myr bins.

Bin start age (Myr)	Bin end age (Myr)	Bin midpoint (Myr)	# of genera
171	166	168.5	0
166	161	163.5	2
161	156	158.5	0
156	151	153.5	12
151	146	148.5	6
146	141	143.5	9
141	136	138.5	10
136	131	133.5	1
131	126	128.5	7
126	121	123.5	11
121	116	118.5	8
116	111	113.5	8
111	106	108.5	7
106	101	103.5	7
101	96	98.5	11
96	91	93.5	5

Bin start age (Myr)	Bin end age (Myr)	Bin midpoint (Myr)	# of genera
91	86	88.5	4
86	81	83.5	16
81	76	78.5	12
76	71	73.5	17
71	66	68.5	21
66	61	63.5	22
61	56	58.5	25
56	51	53.5	11
51	46	48.5	2
46	41	43.5	1
41	36	38.5	1
36	31	33.5	1

Supplementary Table 5. Body mass estimates of multituberculate species. Length and width measurements for lower first molar area (m1 area), skull length measurements (SL), and temporal range data (First Appearance Datum [FAD], Last Appearance Datum [LAD]) were compiled from cited sources. The SL formula is from ref. 39. For species without skull length data, m1 area estimates (ref. 35) were used to predict SL estimates based on a regression analysis (m1-SL; this study).

Species	FAD (Myr)	LAD (Myr)	m1 area (mm²)	SL (mm)	Body mass (kg)	Predictive formula	Data sources
<i>Acheronodon vossae</i>	61	60	2.43	-	0.036	m1-SL	⁵³
<i>Allocosmodon woodi</i>	61	57	2.88	-	0.046	m1-SL	⁵³
<i>Anconodon cochranensis</i>	61	60	3.34	-	0.057	m1-SL	⁵⁴
<i>Arginbaatar dmitrievae</i>	125	100	1.35	-	0.011	m1-SL	⁵⁵
<i>Baiotomeus douglassi</i>	62	61	6.65	-	0.154	m1-SL	⁵⁶
<i>Baiotomeus rhothonion</i>	62	61	1.62	-	0.020	m1-SL	⁵⁷
<i>Barbatodon transylvanicus</i>	71	66	7.37	-	0.178	m1-SL	⁵⁸
<i>Boffius splendidus</i>	62	59	134.69	-	11.808	m1-SL	⁵⁹
<i>Bolodon minor</i>	146	140	0.96	-	0.006	m1-SL	^{60,61}

<i>Bolodon osborni</i>	146	140	1.74	-	0.022	m1-SL	⁶²
<i>Bryceomys fumosus</i>	94	89	1.46	-	0.017	m1-SL	⁶³
<i>Bryceomys hadrosus</i>	94	89	4.28	-	0.081	m1-SL	⁶³
<i>Bryceomys intermedius</i>	100	96	2.16	-	0.030	m1-SL	⁶⁴
<i>Bubodens magnus</i>	69	66	76.80	-	5.248	m1-SL	⁶⁵
<i>Catopsalis alexanderi</i>	66	65	41.92	-	2.190	m1-SL	⁶⁶
<i>Catopsalis calgariensis</i>	61	60	139.73	-	12.451	m1-SL	⁶⁶
<i>Catopsalis fissidens</i>	64	61	82.41	-	5.811	m1-SL	^{67,68}
<i>Catopsalis foliatus</i>	66	64	52.43	-	3.025	m1-SL	⁶⁸
<i>Catopsalis joyneri</i>	66	65	45.11	-	2.435	m1-SL	⁶⁹
<i>Catopsalis waddleae</i>	65	64	152.52	-	14.128	m1-SL	⁷⁰
<i>Catopsbaatar catopsaloides</i>	76	71	19.36	63.00	0.879	SL	^{71,72}
<i>Cedaromys bestia</i>	100	96	3.91	-	0.071	m1-SL	⁶⁴
<i>Cedaromys hutchisoni</i>	79	74	2.29	-	0.033	m1-SL	⁷³
<i>Cedaromys parvus</i>	100	96	2.86	-	0.046	m1-SL	⁶⁴

<i>cf. Paracimexomys perplexus</i>	100	96	1.58	-	0.019	m1-SL	⁶⁴
<i>cf. Paracimexomys robisoni</i>	100	96	1.92	-	0.026	m1-SL	⁶³
<i>Chulsanbaatar vulgaris</i>	84	71	1.53	18.50	0.012	SL	⁷¹
<i>Cimexomys antiquus</i>	84	79	2.37	-	0.035	m1-SL	⁷⁴
<i>Cimexomys arapahoensis</i>	66	65	5.09	-	0.104	m1-SL	⁷⁵
<i>Cimexomys gratus</i>	66	64	5.35	-	0.112	m1-SL	⁷⁶
<i>Cimexomys judithae</i>	79	74	1.97	-	0.027	m1-SL	⁷⁷
<i>Cimexomys minor</i>	69	64	3.10	-	0.051	m1-SL	⁷⁸
<i>Cimolodon electus</i>	84	74	6.63	-	0.153	m1-SL	⁷⁴
<i>Cimolodon foxi</i>	86	74	2.87	-	0.046	m1-SL	⁷⁹
<i>Cimolodon nitidus</i>	86	66	7.40	-	0.179	m1-SL	^{78,80,81}
<i>Cimolodon similis</i>	94	81	5.10	-	0.105	m1-SL	^{74,82}
<i>Cimolodon wardi</i>	84	81	3.87	-	0.070	m1-SL	⁸²
<i>Cimolomys butleria</i>	79	74	4.05	-	0.075	m1-SL	⁷³
<i>Cimolomys clarki</i>	79	74	8.43	-	0.216	m1-SL	⁸³

<i>Cimolomys gracilis</i>	74	66	10.26	-	0.287	m1-SL	78,84
<i>Ctenacodon scindens</i>	155	147	1.58	-	0.019	m1-SL	85,86
<i>Ctenacodon serratus</i>	155	147	1.31	-	0.010	m1-SL	86-88
<i>Dakotamys malcolmi</i>	96	94	2.14	-	0.030	m1-SL	63
<i>Djadochtatherium matthewi</i>	84	81	10.84	-	0.311	m1-SL	89,90
<i>Ectypodus aphronorus</i>	62	60	1.52	-	0.018	m1-SL	54,91
<i>Ectypodus childei</i>	55	49	1.88	-	0.025	m1-SL	92
<i>Ectypodus elaphus</i>	59	58	1.32	-	0.010	m1-SL	93
<i>Ectypodus lovei</i>	43	35	1.56	-	0.019	m1-SL	94,95
<i>Ectypodus musculus</i>	61	56	2.50	-	0.037	m1-SL	96
<i>Ectypodus powelli</i>	61	55	1.90	-	0.025	m1-SL	97
<i>Ectypodus szalayi</i>	62	60	1.76	-	0.022	m1-SL	92
<i>Ectypodus tardus</i>	59	53	1.45	-	0.012	m1-SL	98
<i>Eobaatar clemensi</i>	130	125	1.50	-	0.018	m1-SL	99
<i>Eobaatar magnus</i>	125	100	1.87	-	0.025	m1-SL	55

<i>Essonodon browni</i>	69	66	29.96	-	1.349	m1-SL	⁷⁶
<i>Eucosmodon americanus</i>	65	64	14.56	-	0.476	m1-SL	⁹⁶
<i>Glirodon grandis</i>	155	147	2.21	-	0.031	m1-SL	¹⁰⁰
<i>Guimarotodon leiriensis</i>	155	151	2.85	-	0.045	m1-SL	¹⁰¹
<i>Hainina belgica</i>	66	61	1.63	-	0.020	m1-SL	⁵⁹
<i>Hainina godfriauxi</i>	66	55	5.18	-	0.107	m1-SL	⁵⁹
<i>Hainina pyrenaica</i>	66	65	3.05	-	0.050	m1-SL	¹⁰²
<i>Hainina vianeyae</i>	61	55	4.06	-	0.075	m1-SL	^{102,103}
<i>Heishanobaatar triangulus</i>	125	100	1.86	-	0.024	m1-SL	¹⁰⁴
<i>Iberodon quadrituberculatus</i>	146	140	1.73	-	0.022	m1-SL	¹⁰⁵
<i>Janumys erebos</i>	100	96	0.96	-	0.006	m1-SL	⁶⁴
<i>Kaiparomys cifellii</i>	79	74	4.50	-	0.087	m1-SL	⁷³
<i>Kamptobaatar kuczynskii</i>	84	81	1.98	19.00	0.013	SL	¹⁰⁶
<i>Kimbetohia mziae</i>	66	65	3.92	-	0.072	m1-SL	⁷⁵
<i>Krauseia clemensi</i>	62	60	2.20	-	0.031	m1-SL	^{92,103}

<i>Kryptobaatar dashzevegi</i>	84	71	3.41	32.00	0.083	SL	¹⁰⁶
<i>Kryptobaatar mandahuensis</i>	84	71	3.52	-	0.061	m1-SL	¹⁰⁷
<i>Kuehneodon dietrichi</i>	155	151	2.64	-	0.040	m1-SL	¹⁰⁸
<i>Kuehneodon uniradiculatus</i>	155	151	2.72	-	0.042	m1-SL	¹⁰⁸
<i>Lambdopsalis bulla</i>	59	56	28.00	60.78	0.776	SL	¹⁰⁹
<i>Liaobaatar changi</i>	125	100	4.89	-	0.099	m1-SL	¹¹⁰
<i>Liotomus marshi</i>	61	55	7.13	-	0.170	m1-SL	¹⁰³
<i>Loxaulax valdensis</i>	140	139	3.20	-	0.053	m1-SL	^{55,61,111}
<i>Meketibolodon robustus</i>	155	151	2.97	-	0.048	m1-SL	¹⁰¹
<i>Meniscoessus collomensis</i>	74	66	19.84	-	0.744	m1-SL	¹¹²
<i>Meniscoessus intermedius</i>	86	69	11.13	-	0.323	m1-SL	¹¹³
<i>Meniscoessus major</i>	79	69	15.39	-	0.516	m1-SL	¹¹³
<i>Meniscoessus robustus</i>	74	66	31.92	69.00	1.208	SL	⁸⁶
<i>Meniscoessus seminoensis</i>	69	66	32.21	-	1.497	m1-SL	¹¹⁴
<i>Mesodma ambigua</i>	66	64	3.38	-	0.058	m1-SL	⁹⁶

<i>Mesodma archibaldi</i>	79	74	1.61	-	0.020	m1-SL	⁷³
<i>Mesodma formosa</i>	69	64	2.37	-	0.035	m1-SL	^{78,80,84,115}
<i>Mesodma garfieldensis</i>	66	65	2.73	-	0.042	m1-SL	⁷⁶
<i>Mesodma hensleighi</i>	69	64	1.48	-	0.018	m1-SL	^{80,84,115}
<i>Mesodma minor</i>	79	74	1.26	-	0.009	m1-SL	^{73,79,82}
<i>Mesodma primaeva</i>	79	74	6.00	-	0.132	m1-SL	¹¹⁶
<i>Mesodma pygmaea</i>	62	57	1.02	-	0.006	m1-SL	⁹¹
<i>Mesodma senecta</i>	84	79	3.20	-	0.053	m1-SL	¹¹⁷
<i>Mesodma thompsoni</i>	74	64	3.19	-	0.053	m1-SL	^{80,81,84,115}
<i>Mesodmops dawsonae</i>	55	50	2.59	-	0.039	m1-SL	¹¹⁸
<i>Mesodmops tenuis</i>	56	55	1.98	-	0.027	m1-SL	¹¹⁹
<i>Microcosmodon arcuatus</i>	64	64	2.48	-	0.037	m1-SL	¹²⁰
<i>Microcosmodon conus</i>	59	56	2.78	-	0.044	m1-SL	⁵³
<i>Microcosmodon harleyi</i>	65	64	2.01	-	0.027	m1-SL	¹²¹
<i>Microcosmodon rosei</i>	56	55	2.31	-	0.033	m1-SL	⁹⁷

<i>Mimetodon churchilli</i>	57	56	4.48	-	0.087	m1-SL	⁹⁶
<i>Mimetodon silberlingi</i>	62	57	2.08	-	0.029	m1-SL	¹²²
<i>Nemegtbaatar gobiensis</i>	76	71	4.42	42.00	0.214	SL	⁷¹
<i>Neoliotomus conventus</i>	57	55	23.10	-	0.527	m1-SL	⁹⁷
<i>Neoliotomus ultimus</i>	56	53	23.00	-	0.921	m1-SL	¹³
<i>Neoplagiaulax annae</i>	61	55	3.07	-	0.050	m1-SL	¹⁰³
<i>Neoplagiaulax cimolodontoides</i>	59	58	2.46	-	0.037	m1-SL	⁹³
<i>Neoplagiaulax copei</i>	61	55	4.62	-	0.091	m1-SL	¹⁰³
<i>Neoplagiaulax donaldorum</i>	61	59	3.12	-	0.052	m1-SL	¹²³
<i>Neoplagiaulax eocaenus</i>	61	55	1.77	-	0.023	m1-SL	¹⁰³
<i>Neoplagiaulax grangeri</i>	62	60	4.71	-	0.093	m1-SL	⁵⁴
<i>Neoplagiaulax hazeni</i>	59	56	4.48	-	0.087	m1-SL	⁹⁶
<i>Neoplagiaulax hunteri</i>	62	57	2.98	-	0.048	m1-SL	^{57,122}
<i>Neoplagiaulax macrotomeus</i>	62	61	1.62	-	0.020	m1-SL	¹²⁴
<i>Neoplagiaulax mckennai</i>	59	57	3.51	-	0.061	m1-SL	⁹¹

<i>Neoplagiaulax nicolai</i>	61	55	5.49	-	0.117	m1-SL	¹⁰³
<i>Neoplagiaulax paskapooensis</i>	59	58	2.98	-	0.048	m1-SL	⁹³
<i>Neoplagiaulax serrator</i>	59	58	2.44	-	0.036	m1-SL	⁹³
<i>Nessovbaatar multicosstatus</i>	76	71	2.00	-	0.027	m1-SL	⁹⁰
<i>Nidimys occultus</i>	74	69	5.14	-	0.106	m1-SL	¹²⁵
<i>Paracimexomys magister</i>	84	81	4.48	-	0.087	m1-SL	⁷⁴
<i>Paracimexomys priscus</i>	69	66	3.85	-	0.070	m1-SL	^{80,81}
<i>Paracimexomys propriscus</i>	79	69	2.90	-	0.046	m1-SL	¹²⁵
<i>Parectypodus clemensi</i>	62	61	2.93	-	0.047	m1-SL	¹²⁶
<i>Parectypodus corystes</i>	62	61	2.97	-	0.048	m1-SL	⁵⁷
<i>Parectypodus foxi</i>	69	66	4.17	-	0.078	m1-SL	¹¹⁵
<i>Parectypodus laytoni</i>	58	55	1.36	-	0.011	m1-SL	⁹⁷
<i>Parectypodus lunatus</i>	55	52	2.03	-	0.028	m1-SL	⁹⁸
<i>Parectypodus simpsoni</i>	55	53	2.70	-	0.042	m1-SL	⁹⁸
<i>Parectypodus sinclairi</i>	62	58	1.57	-	0.019	m1-SL	⁵⁴

<i>Parectypodus sylviae</i>	62	60	1.73	-	0.022	m1-SL	126
<i>Parectypodus trovessartianus</i>	62	61	4.65	-	0.092	m1-SL	96
<i>Parikimys carpenteri</i>	69	66	2.89	-	0.046	m1-SL	127
<i>Pentacosmodon pronus</i>	58	56	3.28	-	0.055	m1-SL	53,96
<i>Pinheirodon pygmaeus</i>	146	140	1.23	-	0.009	m1-SL	105
<i>Pinheirodon vastus</i>	146	140	2.07	-	0.028	m1-SL	105
<i>Plagiaulax becklesii</i>	146	140	2.47	-	0.037	m1-SL	60,61
<i>Prionessus lucifer</i>	59	56	12.20	-	0.369	m1-SL	128
<i>Prochetodon cavus</i>	58	56	5.43	-	0.115	m1-SL	129
<i>Prochetodon foxi</i>	59	57	6.83	-	0.160	m1-SL	129
<i>Prochetodon taxus</i>	56	55	6.55	-	0.150	m1-SL	129
<i>Psalodon marshi</i>	155	147	3.42	-	0.059	m1-SL	85,86
<i>Ptilodus fractus</i>	59	56	5.12	-	0.105	m1-SL	130
<i>Ptilodus gnomus</i>	62	59	3.77	-	0.068	m1-SL	57,131
<i>Ptilodus kummae</i>	59	57	5.28	-	0.110	m1-SL	122

<i>Ptilodus mediaevus</i>	64	56	6.93	-	0.163	m1-SL	126,132
<i>Ptilodus montanus</i>	62	57	5.98	38.97	0.165	SL	54,132
<i>Ptilodus wyomingensis</i>	62	57	6.44	-	0.147	m1-SL	96
<i>Sinobaatar fuxinensis</i>	125	100	2.71	-	0.042	m1-SL	110
<i>Sinobaatar xiei</i>	125	100	2.40	-	0.035	m1-SL	110
<i>Sloanbaatar mirabilis</i>	84	81	1.70	22.30	0.023	SL	106
<i>Stygmymys jepseni</i>	62	61	4.37	-	0.084	m1-SL	54
<i>Stygmymys kuszmauli</i>	66	64	8.30	-	0.212	m1-SL	76
<i>Taeniolabis lamberti</i>	65	64	128.00	-	10.971	m1-SL	133
<i>Taeniolabis taoensis</i>	64	64	205.92	160.00	22.697	SL	133
<i>Uzbekbaatar wardi</i>	91	86	2.20	-	0.031	m1-SL	134
<i>Xyronomys robinsoni</i>	66	65	1.87	-	0.025	m1-SL	75

Supplementary Table 6. OPC and OPCR results for carnivorans and rodents in Evans et al.¹⁸.

Species	OPC (l)	OPCR (l)	OPC (u)	OPCR (u)
<i>Acinonyx jubatus</i>	40	36.5	53	52.625
<i>Ailuropoda melanoleuca</i>	257	266.625	342	326.25
<i>Ailurus fulgens</i>	195	194.625	270	269.875
<i>Alopex lagopus</i>	95	96.75	174	154.25
<i>Canis aureus</i>	125	132.875	185	172.625
<i>Canis lupus</i>	97	97.375	137	135.25
<i>Crocuta crocuta</i>	61	55.875	72	76.125
<i>Felis silvestris</i>	37	44.625	55	62.25
<i>Galerella sanguinea</i>	110	107.125	143	144.25
<i>Genetta genetta</i>	140	139.5	144	143.875
<i>Gulo gulo</i>	54	54.25	124	122.5
<i>Herpestes ichneumon</i>	101	101.125	149	144.25
<i>Lutra lutra</i>	91	89.875	150	152.875
<i>Lynx lynx</i>	40	44.75	58	62.25
<i>Martes foina</i>	79	72.5	145	143.75
<i>Martes martes</i>	77	79.875	142	138.125
<i>Meles meles</i>	121	118.875	195	194.625
<i>Mustela erminea</i>	39	43.5	76	76.875
<i>Mustela eversmannii</i>	48	46.75	91	83.5
<i>Mustela lutreola</i>	68	71.25	126	119.25
<i>Mustela nivalis</i>	54	54.5	90	89.875
<i>Mustela putorius</i>	55	55.875	99	99.625
<i>Otocyon megalotis</i>	162	163.625	184	176.5
<i>Panthera leo</i>	37	41.125	53	57.375
<i>Paradoxurus leucomystax</i>	115	121.75	115	114.25

<i>Procyon lotor</i>	153	154.75	188	180.375
<i>Ursus americanus</i>	192	194.125	150	164.5
<i>Ursus arctos</i>	179	185.625	170	172.5
<i>Ursus maritimus</i>	201	189.75	135	137.875
<i>Viverra zibetha</i>	162	161.375	206	203
<i>Vormela peregusna</i>	56	56.75	113	113.25
<i>Vulpes vulpes</i>	119	117.625	184	173.25
<i>Aethomys hindei</i>	225	222.25	231	241.625
<i>Akodon serrensis</i>	113	114.75	129	128
<i>Anisomys imitator</i>	231	235.125	280	280.875
<i>Apodemus agrarius</i>	173	176.375	173	176.125
<i>Arvicanthis niloticus</i>	150	153.625	182	185.875
<i>Berylmys bowersi</i>	193	187.875	209	204.125
<i>Crateromys schadenbergi</i>	230	233.375	285	283.75
<i>Dasymys</i> sp.	233	233.875	309	305.25
<i>Geoxus valdivianus</i>	184	175.625	191	184.5
<i>Grammomys dolichurus</i>	229	228.5	265	265
<i>Grammomys rutilans</i>	220	238.25	273	282.625
<i>Holochilus brasiliensis</i>	287	281.75	285	275.875
<i>Hybomys univittatus</i>	220	215.5	233	234.625
<i>Hydromys chrysogaster</i>	122	122.125	126	133.125
<i>Hylomyscus stella</i>	190	189.375	177	180.25
<i>Hyomys goliath</i>	277	269.375	309	303.625
<i>Ichthyomys stolzmanni</i>	163	159.875	182	184.125
<i>Lemniscomys striatus</i>	167	166.75	218	222.625
<i>Leopoldamys sabanus</i>	217	227.875	229	232.25
<i>Leptomys elegans</i>	149	147.875	134	132.375
<i>Lophuromys medicaudatus</i>	167	165.25	256	259.875
<i>Malacomys</i> sp.	180	187.25	206	207.625
<i>Mallomys rothschildi</i>	241	241.5	259	259.375

<i>Mastomys natalensis</i>	152	155.375	167	179.875
<i>Melomys levipes</i>	177	182.375	189	196
<i>Micromys minutus</i>	164	166.5	162	164.75
<i>Mus musculus</i>	167	173.375	181	176.625
<i>Nectomys squamipes</i>	226	220.625	253	260.875
<i>Nesokia indica</i>	191	189.125	237	236.125
<i>Notomys mitchellii</i>	181	180.125	203	206.25
<i>Oenomys hypoxanthus</i>	187	198.25	190	192.375
<i>Otomys denti</i>	146	140	197	189.375
<i>Otomys irroratus</i>	133	128.75	167	172.75
<i>Oxymycterus</i> sp.	153	158.875	130	140
<i>Parotomys littledalei</i>	173	176.75	212	222.75
<i>Pelomys campanae</i>	189	191.875	196	202.5
<i>Peromyscus maniculatus</i>	200	190.5	201	204.875
<i>Phyllotis</i> sp	214	215.875	181	183.875
<i>Pogonomys</i> sp	246	247.5	293	279.75
<i>Praomys jacksoni</i>	177	178.375	162	162.25
<i>Rattus leucopus</i>	239	246.125	236	236.75
<i>Reithrodon auritus</i>	222	232	247	250.5
<i>Reithrodontomys mexicanus</i>	256	252.25	276	257.5
<i>Rhabdomys pumilio</i>	180	177.625	185	188.25
<i>Sigmodon hispidus</i>	267	258	276	293.875
<i>Stochomys longicaudatus</i>	265	269	266	267.25
<i>Sundamys muelleri</i>	203	212.625	245	232.625
<i>Uromys caudimaculatus</i>	184	202.75	197	207
<i>Zygodontomys</i> sp.	178	175.625	202	189.625

Supplementary Table 7. Comparisons of OPC results for two multituberculate tooth row casts (*Meniscoessus robustus* UCMP 107405, *Parikimys carpenteri* DMNH 52224) for three 3D scanners. Relative standard errors for the two casts were 0.98% and 3.06% respectively.

Species	Scanner	OPCR
<i>M. robustus</i>	MicroCT	168.375
<i>M. robustus</i>	Hawk	164.25
<i>M. robustus</i>	Laser Design	163
<i>P. carpenteri</i>	MicroCT	144.875
<i>P. carpenteri</i>	Hawk	134.875
<i>P. carpenteri</i>	Laser Design	130.75

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