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Reconstructing the archosaur radiation using a Middle Triassic archosauriform tooth assemblage from Tanzania

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Following the Permo-Triassic mass extinction, Archosauriformes – the clade that includes crocodylians, birds, and their extinct relatives – rapidly diversified into numerous distinct lineages, became distributed globally, and, by the Late Triassic, filled a wide array of resource zones. Current scenarios of archosauriform evolution are ambiguous with respect to whether their taxonomic diversification in the Early–Middle Triassic coincided with the initial evolution of dietary specializations that were present by the Late Triassic, or if their ecological disparity arose sometime after lineage diversification. Late Triassic archosauriform dietary specialization is recorded by morphological divergence from the plesiomorphic archosauriform tooth condition (laterally-compressed crowns with serrated carinae and a generally triangular lateral profile). Unfortunately, the roots of this diversification are poorly documented, with few known Early–Middle Triassic tooth assemblages, limiting characterizations of morphological diversity during this critical, early period in archosaur evolution. Recent fieldwork (2007–2017) in the Middle Triassic Manda Beds of the Ruhuhu Basin, Tanzania, recovered a tooth assemblage that provides a window into this poorly sampled interval. To investigate the taxonomic composition of that collection, we built a dataset of continuous quantitative and discrete morphological characters based on *in situ* teeth of known taxonomic status (e.g., *Nundasuchus*, *Parringtonia*: N = 65) and a sample of isolated teeth (N = 31). Using crown heights from known taxa to predict tooth base ratio (= base length/base width), we created a quantitative morphospace for the tooth assemblage. The majority of isolated, unassigned teeth fall within a region of morphospace shared by several Manda taxa (e.g., *Nundasuchus*, *Parringtonia*); two isolated teeth fall exclusively within a 'Pallisteria' morphospace. A non-metric multidimensional scaling ordination (N = 67) of 11 binary characters reduced overlap between species. The majority of the isolated teeth from the Manda assemblage fall within the *Nundasuchus* morphospace. This indicates these teeth

are plesiomorphic for archosauriforms as *Nundasuchus* exhibits the predicted plesiomorphic condition of archosauriform teeth. Our model shows that even the conservative tooth morphologies of archosauriforms can be differentiated and assigned to species/genus level, rendering the model useful for identifying isolated teeth. The large overlap in tooth shape among the species present and their overall similarity indicates that dietary specialization lagged behind species diversification in archosauriforms from the Manda Beds, a pattern predicted by Simpson's 'adaptive zones' model. Although applied to a single geographic region, our methods offer a promising means to reconstruct ecological radiations and are readily transferable across a broad range of vertebrate taxa throughout Earth history.

1 Reconstructing the Archosaur Radiation using a Middle

2 Triassic Archosauriform Tooth Assemblage from Tanzania

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42 region, our methods offer a promising means to reconstruct ecological radiations and are readily
43 transferable across a broad range of vertebrate taxa throughout Earth history.

44 **Introduction**

45 Adaptive radiations, or evolutionary diversifications, play a critical role in the history of life as
46 clades speciate and fill new ecological roles over geologically rapid time intervals (Simpson,
47 1944; Schlüter, 1996). Although there are examples of adaptive radiations that are not speciose
48 (e.g. Darwin's finches), or adaptively disparate (e.g. crotaphytine and oplurine iguanids), such a
49 framework is still useful for structuring macroevolutionary questions and explaining present (and
50 past) biological diversity (Gavrilets and Losos, 2009). Adaptive radiations and the shifts in
51 evolutionary rates associated with them are among the most studied aspects of evolutionary
52 biology (e.g. Stanley, 1979; Losos and Miles, 1994, 2002; Gavrilets and Losos, 2009; Revell et
53 al., 2018; Slater and Friscia, 2019). However, empirical uncertainties remain regarding many of
54 the properties of adaptive radiations (Gavrilets and Losos, 2009; Slater and Friscia, 2019), with
55 the relative timings of lineage diversification and ecological disparity during adaptive radiations
56 being one such problem. Does lineage diversification come first, followed by specialization and
57 evolution within an 'adaptive zone' (Simpson, 1944, 1953) or does ecological specialization
58 drive lineage diversification simultaneously (Schlüter, 1996)? In the former case species fill the
59 same resource zones using similar, ancestral morphological structures (e.g. identical tooth
60 morphologies), whereas in the latter each species would be expected to have a unique, derived
61 morphology for its resource zone at the start of the radiation (for an empirical example, see
62 Slater and Friscia, 2019). Determining which of these competing hypotheses operated in a
63 particular case requires us to reconstruct an evolutionary radiation where a species-poor,
64 adaptively **constrained** clade diversifies into a species-rich, adaptively disparate clade.

65 One such radiation occurred during the Triassic Period, following the Permo-Triassic
66 mass extinction (PTME: Raup, 1979; Erwin, 1994; Chen and Benton, 2012; Benton and Newell,

67 2014) as archosauriforms recovered, rapidly diversified, and spread across Pangea to dominate
68 terrestrial ecosystems for the next 150 million years (Nesbitt, 2011; Ezcurra and Butler, 2018). In
69 addition to Archosauriformes being a speciose and disparate radiation, they also provide an
70 opportunity to test adaptive radiations at a higher phylogenetic level. Lineage diversification of
71 archosaurs was rapid after the PTME, and by the Middle Triassic many non-archosaurian
72 archosauriform and crown archosaur clades had appeared (Ezcurra, 2016; Foth et al., 2016;
73 Ezcurra and Butler, 2018). By the Late Triassic archosaurs filled a wide variety of ecological
74 roles, from top predators to large herbivores, and were represented in terrestrial, freshwater, and
75 even marine ecosystems (e.g. Li et al., 2006; Butler et al., 2019). If lineage diversification occurs
76 first, followed by subsequent ecological disparity, we would expect Middle Triassic
77 archosauriforms from across the tree to present a limited range of ecologies. The question that
78 arises is how best to measure ecological disparity? Ecological disparity covers a variety of
79 physiological, behavioral, and morphological traits but the nature of the fossil record limits its
80 measure primarily to morphology. Previous work has used cranial morphology as a measure of
81 disparity (e.g. Foth et al., 2016); however complete, or even partial, skulls are rare for Early–
82 Middle Triassic archosauriforms. Therefore, an alternative morphological system to approximate
83 ecological disparity is needed. In this study we use teeth as an indicator of ecological disparity
84 because they have relatively high preservation potential (e.g. Turner-Walker, 2008) and offer a
85 direct link to ecology through diet (Lucas, 1979; Dessem, 1985; Scallon and Shine, 1988;
86 Sander, 1997; Linde, Palmer & Gómez-Zurita, 2004; Santana, Strait & Dumont, 2011;
87 Zahradnicek et al., 2014). We consider diet as the aspect of ecology of interest for both the
88 relative ease of inference from morphology alone and the use of diet in previous studies of
89 evolutionary radiations (e.g. Slater and Frisia, 2019).

90 Although tooth assemblages are rare in Middle Triassic terrestrial settings, recent
91 fieldwork (2007, 2008, 2012, 2015, 2017) in the Manda Beds of the Ruhuhu Basin, Tanzania
92 (Sidor and Nesbitt, 2017), has revealed a rich assemblage of archosauriforms known from
93 postcrania and partial crania, including teeth (e.g. Nesbitt et al., 2010; 2014; Smith et al., 2018).
94 Specifically, these teeth come from the middle and upper Lifua Member bone accumulations
95 (Smith et al., 2018), which are thought to be Anisian in age (Rubidge, 2005) but may be as
96 young as early Carnian (Ottone et al., 2014; Marsicano et al., 2016; Wynd et al., 2018; Peecook
97 et al., 2018). If the Anisian age is correct, then this is one of the oldest, diverse archosaur faunas
98 known that is also represented by specimens from historical collections (e.g. Butler et al., 2009;
99 2017; Nesbitt et al., 2010; 2013; 2014; 2017; Barrett, Nesbitt & Peecook, 2015). Using a
100 combination of information from these new and historical collections, we quantify tooth
101 disparity in this earliest part of the archosaur radiation to generate a morphospace visualization.
102 From this we can assign isolated teeth to specific taxon, visualize inter- and intraspecific
103 variation as well as intra-individual variation, and use this variation as a window into the
104 ecological disparity of the archosauriforms within the Lifua Member assemblage. To achieve
105 these goals we use a combination of *in situ* teeth from jaw elements assignable to particular
106 species (Figure 1), and isolated teeth attributable to Archosauriformes (Figure 2). Of particular
107 interest is whether the isolated teeth fall within, or expand, the region of morphospace occupied
108 by the described Manda Beds taxa.

109 *Institutional Abbreviations* – **NHMUK**, Natural History Museum, London, U.K.; **NMT**,
110 National Museum of Tanzania, Dar es Salaam, Tanzania.

111

112 **Materials & Methods**

113 The 31 isolated teeth included in this study were ~~all~~ collected from surface accumulations
114 ~~of vertebrate fossils~~ during fieldwork in 2007, 2008, 2012, 2015, and 2017 from the Manda Beds
115 of the Ruhuhu Basin by a multi-institutional team (Sidor and Nesbitt, 2017). All of the isolated
116 teeth included in this study are currently housed at Virginia Tech Department of Geosciences and
117 will be permanently reposed and managed in the National Museum of Tanzania. In addition to
118 these isolated teeth (seven of which were referred to *Nundasuchus*: see Nesbitt et al., 2014), we
119 also included teeth from within the tooth-bearing elements of five taxonomically distinct
120 archosauriforms from the Manda Beds: *Nundasuchus* (NMT RB48), *Parringtonia* (NMT
121 RB426), *Asilisaurus* (NMT RB837), 'Pallisteria' (NHMUK PV R36620), and one currently
122 undescribed pseudosuchian that we refer to ~~–~~ by its specimen number (NMT RB187). We assign
123 the isolated teeth to Archosauriformes on the basis of their serration morphology (Nesbitt, 2011)
124 as well as their general ziphodont construction, including lateral compression (e.g. Godefroit and
125 Cuny, 1997).

126 To quantify tooth shape, linear measurements (total crown height, base width, and fore-
127 aft base length) and denticle counts were made following the protocol in Smith, Vann & Dodson
128 (2005), although due to the smaller size of the teeth in our study, we used 1 mm denticle
129 densities, rather than 5 mm densities (Supplemental Data S1). All statistical analyses were
130 performed in R (v 3.1.2) and the RStudio console (v 1.1.383). All graphs of quantitative data
131 were made using the R package "ggplot2" (Wickham, 2009). To capture tooth disparity (from
132 log-transformed linear measurements) we used sum of variances with 95% predictive intervals
133 following the methodology of Larson, Brown & Evans (2016). We chose to use sum of variances
134 as our measure of disparity due to its prevalence in the literature and its robustness when
135 working with small sample sizes. Sample size varied from 2–14 teeth, however, sample size does

136 not significantly affect the sum of variance analysis (Ciampaglio, Kemp, & McShea, 2001). In
137 *Asilisaurus* the 14 teeth largely contributed base and width measurements because only three
138 teeth were complete enough to measure crown height. We constructed a linear model in R
139 predicting the variable of tooth base shape (ratio of mesiodistal length over labiolingual width)
140 by the variables of total crown height and species-level assignment such that base shape = total
141 crown height * species assignment. The effects of each species on predicting tooth base shape
142 were elucidated using the R package “lsmeans” (Lenth, 2016) using a pairwise comparison in the
143 model by taxon. We plotted the teeth of known taxonomic affinity using ggplot2 (Wickham,
144 2009) to produce a base morphospace into which we plotted results from the isolated teeth for
145 comparison.

146 Simple quantitative measurements only capture the overall shape of the teeth, and all of
147 the teeth in the study resemble the hypothetical ancestral archosauriform tooth (serrated,
148 recurved, and laterally compressed: Nesbitt, 2011). In order to more fully capture and describe
149 the subtle variation of these teeth, a method of capturing discrete variation is needed. Non-metric
150 multidimensional scaling (NMDS) is an ordination method that visualizes variation that can
151 incorporate discrete qualitative features. We created a set of 11 binary characters for scoring
152 isolated and *in situ* teeth for NMDS (Table 1, Figure 3, Supplemental Data S2). All characters
153 except one are new to this analysis (trait 6, dental caudae = shallow grooves extending from
154 between two adjacent denticles present/absent is taken from Abler [1992]). None of the traits
155 used in this study have been used in phylogenetic analyses of archosauriforms, in order to avoid
156 circular reasoning when comparing our ecological signal to taxonomic and clade identity. The
157 NMDS analysis was conducted in PAST (Hammer, Harper & Ryan, 2001) with a Bray-Curtis
158 transformation. We ran an additional NMDS analysis in PAST using average taxon and

159 morphotype scores where traits were scored for each taxon with >50% agreement of *in situ* teeth.

160 Traits for which <50% of the specimens in the taxon or morphotype were scored as unknown

161 (“?”).

162

163 **Isolated Tooth Descriptions**

164 *Morphotype A*: These teeth (Figure 2A) are generally triangular in outline in lateral view and

165 most are recurved (the point of the crown is distal to the distal-most extent of the crown base)

166 although the remainder have crown tips that are level with the distal-most extent of the crown

167 base. The labial and lingual sides of the crown lack ridges (i.e. no fluting), and the labial side of

168 the crown exhibits greater convexity than the lingual side. The mesial denticle series terminates

169 more apically along the crown margin than the distal series, which continues along the entire

170 height of the crown though both start at the tip of the crown. The mesial denticle series is also

171 offset from the mesial-distal long axis of the crown base, deflecting to the lingual side toward the

172 crown base. The denticle densities range from 2–5 per mm. Denticle caudae (Abler, 1992),

173 which are shallow grooves extending from between two adjacent denticles, are often present and

174 directed parallel to the denticles. These denticle caudae are most easily viewed in mesial or distal

175 view (Figure 3F).

176 In general, Morphotype A teeth strongly resemble both *in situ* and isolated teeth of

177 *Nundasuchus* (Figure 1; Nesbitt et al., 2014). Particularly important features are the presence of

178 denticle caudae, an unequal labial-lingual curvature, and the more apical termination of the

179 mesial denticle series relative to the distal denticle series. Also like *Nundasuchus*, teeth of

180 Morphotype A exhibit a mix of states in the changing curvature of the mesial crown edge in

181 lateral view, with some teeth gradually changing angles and others exhibiting an abrupt shift in

182 angle. The *in situ* teeth of *Nundasuchus* can exhibit either state depending on the proximity of
183 the mesial edge of the crown to the distal edge of the preceding tooth. Though this combination
184 of traits is only found in *Nundasuchus* in the Manda Beds fauna, archosauriforms from the
185 Middle and Upper Triassic elsewhere (e.g. de Oliveira and Pinheiro, 2017; Schoch et al., 2018).

186 *Morphotype B*: These teeth (Figure 2B) are triangular in shape in lateral view and are
187 occasionally recurved, although in most the apical tip of the crown is approximately level with
188 the distal-most end of the crown base. Morphotype B tooth crowns lack fluting and, in contrast to
189 Morphotype A, the labial and lingual curvatures are equal. None of the teeth are bulbous (no
190 labiolingual measurements are greater than crown base width). In the majority of Morphotype B
191 teeth the mesial margin of the crown possesses a single point where the angle of the mesial
192 carina changes abruptly. As in Morphotype A teeth, the mesial series of denticles in Morphotype
193 B teeth terminates on the crown further apically than the distal series, which often terminates at
194 the crown base. However, the mesial row of denticles is in line with the mesial-distal long axis of
195 the crown base. The denticle densities range from 3–8 per mm. Denticle caudae are present on
196 some of the teeth and are directed parallel to the denticles. Although these teeth bear a strong
197 resemblance to Morphotype A, they can be differentiated by their equal labial and lingual
198 curvatures. Morphotype B teeth are similar to some of the *in situ* and isolated *Nundasuchus* teeth
199 (Figure 1,2; Nesbitt et al., 2014).

200 *Morphotype C*: This morphotype (Figure 2C) is represented by a single tooth in our assemblage,
201 NMT RB831. The overall shape is tall, near conical, and recurved. The crown lacks fluting and
202 the labial curvature is greater than the lingual curvature. Although its labial-lingual curvature is
203 unequal, the mesial denticle series is positioned along the midline of the mesial-distal long axis.
204 The orientation of the mesial edge of the tooth changes gradually, forming a long, continuous

205 curve. The tooth is not bulbous. Denticle densities range from 2–4 per mm, and no denticle
206 caudae are present. There is no variation in either the shape or size of the denticles between the
207 mesial and distal series or along the length of the crown. Unlike either Morphotypes A or B, the
208 mesial series of denticles in Morphotype C ends at approximately the same level on the crown as
209 the distal series, just above the crown base.

210 In general size and shape, as well as in many of its discrete features, the Morphotype C
211 tooth is similar to the teeth of '*Pallisteria*' based on our observations. The teeth of the latter
212 taxon are large, conical, recurved, and possess unequal labial-lingual curvature. The denticle
213 density is low (< 3 per mm) in the middle part of the tooth crown and denticles show little
214 variation in shape or size. Unfortunately, none of the '*Pallisteria*' teeth could be scored for Trait
215 7 (termination height of the mesial denticle series; Table 1) due to poor preservation of the
216 denticles, which otherwise differentiates Morphotype C teeth from morphotypes A and B. If
217 Morphotype C is similar to, or is, '*Pallisteria*', then subsequent '*Pallisteria*' tooth discoveries
218 should be expected to have sub-equally extending mesial and distal denticle rows.

219

220 ***In situ* Tooth Descriptions**

221 *Nundasuchus*: We included a total of 13 *Nundasuchus* teeth, six *in situ* and seven isolated,
222 originally described in Nesbitt et al. (2014). The teeth range in height from 5.6 to 22 mm with
223 denticle densities from 2–5 per mm. All the teeth are labio-lingually compressed and are serrated
224 on both mesial and distal margins. Only one tooth possesses a recurved tip that extends past the
225 distal-most end on the tooth base. Most teeth are smooth on the sides with a single exception
226 exhibiting fluting. All of the teeth possess: unequal labial-lingual curvatures, a mesial row of
227 denticles that terminates higher on the tooth crown than the distal row of denticles, and a mesial

228 carina that is offset from the midline. Only two of the teeth possess dental caudae and one tooth
229 is bulbous. In some teeth the mesial and distal denticle rows differ in size and/or in shape. About
230 half the teeth have a distinct point on the mesial margin where the angle of the edge changes
231 abruptly. For the *in situ* teeth this seems to be related to how close the tooth is to the preceding
232 socket, with the closer the distance being associated with an abrupt angle shift point.

233 *Asilisaurus*: We included 14 *in situ* teeth though only three of these included more than the very
234 base of the tooth. These three ranged in height from 1.6 to 2.9 mm and had a denticle density of
235 ~8 per mm. The teeth are closely packed, ankylosed to the sockets, and peg-like in shape
236 (Nesbitt et al., 2010). All of the teeth have: smooth sides, equal labial-lingual curvature, and
237 subeven mesial and distal row of denticles. None of the *Asilisaurus* teeth possess dental caudae
238 and the mesial edge of the teeth changes angles gradually.

239 *Parringtonia*: Of the 14 teeth in the study, 12 were *in situ* and the other two larger, isolated teeth.
240 The teeth range in size from 2.5–21.6 mm, though the tallest *in situ* tooth is 8.3 mm, and the
241 denticle densities vary from 5–15 per mm. Most of the *Parringtonia* teeth lacked crown tips,
242 though the two complete teeth are not recurved. All of the teeth are labio-lingually compressed
243 and possess fluting and a mesial carina along the midline. The mesial and distal denticle series of
244 all the teeth remain constant in both shape and size, though the mesial denticle series terminates
245 higher on the crown than the distal series. In all the teeth the mesial edge angle changes
246 gradually.

247 NMT RB187: All 13 teeth of the teeth included from NMT RB187 are *in situ*. The labio-
248 lingually compressed teeth range from 5.3–13.4 mm tall with denticle densities of 8–14 per mm.
249 All of the teeth are recurved, fluted, and lack dental caudae. The mesial edge of the teeth changes
250 gradually and follows the mesial-distal long axis. In teeth with preserved crown tips the shape of

251 the denticles remains constant. Of all the taxa included here, NMT RB187 exhibits the greatest
252 degree of recurvedness.

253 ‘*Pallisteria*’: We included 11 *in situ* teeth from the left and right maxillae of ‘*Pallisteria*’. These
254 teeth are the largest of all the taxa, ranging from 36.1–70.3 mm, and have the lowest density,
255 from 2–3 per mm. All except one are recurved and all have smooth crowns and lack dental
256 caudae. Most of the teeth have uneven labial-lingual curvature and a mesial edge that changes
257 angles gradually. The mesial carina is offset from the mesiodistal long axis in most the teeth and
258 the denticles remains constant in shape and size along the height of the crown.

259

260 **Results**

261 For our linear model we predicted the tooth base shape (ratio of labiolingual base width
262 to mesiodistal base length) using the total apicobasal crown height and the taxonomic affinity of
263 the tooth (base ~ tch + taxon) with the lm() command in base R (Table 2). We found that tooth
264 height was not a significant predictor of base shape ($p = 0.0933$). We used the R package
265 “lsmeans” to further investigate the differences between the species’ tooth shape (Table 3). From
266 this metric NMT RB187 has a significantly higher base shape ratio than all other taxa except
267 *Parringtonia* ($p = 0.3788$).

268 The sum of variances analysis (Figure 4) included all known Manda Beds archosauriform
269 taxa with associated dentition and two of the three morphotypes, as only a single tooth of
270 Morphotype C is present in our assemblage. These variances provide a quantification of
271 intraspecific variation in tooth size and shape, and allow for an equal interspecific comparison.
272 Mean variances ranged from a low of 0.02 log units in ‘*Pallisteria*’, two large isolated teeth of
273 *Parringtonia*, and Morphotype B, to a high of 0.145 log units in Morphotype A (Figure 4).

274 More useful for visualizing variation than the linear model and lsmeans contrasts are
275 morphospace plots of the teeth from our generically determinate specimens, with the isolated,
276 unidentified teeth added for comparison. There is much overlap in morphospace occupancy,
277 particularly on the left side (shorter height) portion of the graph, although '*Pallisteria*' occupies
278 its own section of morphospace in taller crown heights (Figure 5). Teeth towards the bottom of
279 the morphospace (lower base ratio) are more rounded and cone-like, whereas those with higher
280 base ratios are more laterally compressed. With size alone two of the Morphotype A teeth fall in
281 '*Pallisteria*' morphospace and the Morphotype C tooth with *Nundasuchus* morphospace contrary
282 to the discrete descriptive predictions. The relationship between base width and mesiodistal base
283 length provides little more distinction of the taxa included, and the impact of crown size is still
284 evident (Figure 6). In general the ratio of base mesiodistal length and labiolingual width follows
285 a linear trend controlled by size.

286 A total of 21 isolated teeth and 46 *in situ* teeth of known affinity were complete enough
287 to be scored for the NMDS analysis. Convex hulls are more differentiated than in the quantitative
288 morphospace, with almost no overlap of *Nundasuchus* with either NMT RB187 or *Parringtonia*
289 (Figure 7). Overlap of NMT RB187 and *Parringtonia* remains, but most of the isolated teeth fall
290 exclusively within or adjacent to the zone of *Nundasuchus* and '*Pallisteria*' (Figure 7). The high
291 degree of overlap between *Parringtonia* and NMT RB187 likely reflects their often-shared
292 feature of having parallel ridges (fluting) along the labial and lingual sides of the tooth crown.
293 The only other tooth in the study with fluting is a single example referred to *Nundasuchus*. The
294 use of taxa and morphotype 'averages' in traits reveals similar groupings to the complete dataset,
295 with average morphotype scores between those of known taxa (Figure 8).

296

297 **Discussion**

298 We present the first quantitative description of a Middle Triassic archosauriform tooth
299 assemblage, which reveals substantial conservation of tooth morphology at the beginning of the
300 archosaur radiation. Intraspecific variation appears to be as great, if not greater, than interspecific
301 variation. Morphotype A displays the greatest variance in tooth size in the sample, although
302 *Nundasuchus* has a very similar sum of variance structure (Figure 4). Driving at least part of the
303 pattern we see in our disparity analysis is whether more than a single individual of a given taxon
304 is included in our study. For example, NMT RB187, ‘*Pallisteria*’, and *Parringtonia* all display
305 low disparity, but our sample includes only elements from a single individual of each taxon,
306 whereas the *Nundasuchus* sample includes *in situ* teeth from one lower jaw (the holotype
307 specimen) and associated isolated teeth assigned to the holotype (Figure 4). Although two of the
308 isolated teeth from our assemblage fall exclusively within the ‘*Pallisteria*’ quantitative
309 morphospace, most of the isolated teeth fall within a zone of overlap between *Nundasuchus*,
310 NMT RB187, and *Parringtonia* (Figure 5). Much of this quantitative variation reflects body size
311 (Figure 6). *Nundasuchus* and ‘*Pallisteria*’ are much larger than the other taxa, which helps to
312 differentiate their morphospace from that of smaller-bodied taxa. *Asilisaurus* is the smallest
313 taxon in our sample, but there is postcranial evidence of a larger silesaurid in the Lifua
314 assemblage (possibly a very large individual of *Asilisaurus*; Barrett, Nesbitt, & Peecook, 2015)
315 that would be comparable in size to *Nundasuchus* and ‘*Pallisteria*’. Recovery of teeth from
316 silesaurid individuals of this larger size might reduce some of the differentiation between them,
317 *Nundasuchus*, and ‘*Pallisteria*’ though we would still expect silesaurid teeth to be smaller
318 relative to the same body size.

319 The NMDS ordination improves the differentiation of taxa, with *Asilisaurus* and the
320 large-bodied predator '*Pallisteria*' more clearly separated from the still overlapping undescribed
321 pseudosuchian, and *Parringtonia* and *Nundasuchus* exhibiting wide variation in morphospace
322 overall, bridging the space between all taxa, and overlapping a substantial part of '*Pallisteria*'
323 morphospace (Figure 7). These results identify two general areas of morphospace, one shared by
324 the undescribed pseudosuchian and *Parringtonia* and the other by *Nundasuchus* and
325 '*Pallisteria*'. The teeth of *Parringtonia* and the undescribed pseudosuchian share several
326 features, notably presence of fluting, a mesial carina along the midline tooth axis, and a high
327 denticle density (≥ 3 per mm). By contrast, *Nundasuchus* and '*Pallisteria*' teeth lack fluting,
328 possess an offset mesial carina, unequal labial/lingual curvature, and have a low denticle density
329 (< 3 per mm). This result is further supported when the average or typical score of each taxon is
330 used, with NMT RB187, *Asilisaurus*, and *Parringtonia* clustering together versus *Nundasuchus*
331 and '*Pallisteria*' on the other side of morphospace (Figure 8). Given that many of the isolated
332 teeth resemble those of *Nundasuchus*, it is not surprising that most of the isolated teeth fall
333 within the convex hull defined by *Nundasuchus* (Figure 7). We cannot, however, definitely
334 assign these teeth to *Nundasuchus* due to the overlap in discrete characters among our included
335 taxa.

336 Our results using both methods demonstrate that many of the isolated teeth resemble
337 those from currently recognized taxa. However, several teeth fall outside of the morphospace
338 defined by known taxa and could indicate either intraspecific variation (due to heterodonty or
339 ontogeny) or could represent other, as yet unsampled, taxa. Our methodologies are flexible and
340 the datasets can incorporate additional specimens as they are excavated, so these approaches

341 could be applied to other tooth assemblages throughout the Triassic across a broad range of
342 spatial, temporal, and taxonomic scales.

343 **Ecological Differentiation.** There are some hints of dietary separation between large-
344 and small-bodied archosaurs based on minor changes in tooth morphology and consideration of
345 body size. However, our results, which show high degrees of overlap in tooth morphology
346 suggest that ecological differentiation, at least in diet, appears to lag behind lineage
347 diversification, at least with respect to Manda archosauriforms. Four of the five recognized taxa
348 included here possess ziphodont dentitions (=labiolingual narrow crown [labiolingual width <
349 60% of mesiodistal length], recurved, typically serrated carinae, and no constriction at the cervix
350 *sensu* Hendrickx, Mateus & Araújo, 2015) indicative of a carnivorous diet. Only *Asilisaurus*
351 differs in possessing a conodont dentition (=conical crowns with small denticles or no denticles,
352 and typically fluted *sensu* Hendrickx, Mateus & Araújo, 2015). Conodonty is present in
353 spinosaurids, many crocodylians, marine reptiles, and pterosaurs (Hendrickx, Mateus & Araújo,
354 2015) and has been linked to piscivory. Following this criterion *Asilisaurus* would be categorized
355 as a potential piscivore. However, dietary reconstructions of *Silesaurus opolensis*, another
356 silesaurid possessing similar dentition to *Asilisaurus*, have been herbivorous or omnivorous
357 based upon dental microwear (Kubo and Kubo, 2014) or insectivorous based upon coprolites
358 (Qvarnström et al., 2019). Thus, in the Manda Beds tooth assemblage there are two large-bodied
359 carnivores (*Nundasuchus* and ‘*Pallisteria*’), two small-bodied carnivores (*Parringtonia* and an
360 undescribed pseudosuchian), and one small-bodied, non-carnivore (*Asilisaurus*). The Middle
361 Triassic Manda Beds may, therefore, be capturing the beginning of the ‘Explosive Phase’ of
362 Simpson’s (1944) theoretical model as lineages split and begin to move towards new adaptive
363 zones. Further tooth assemblages will need to be evaluated to see if this is a broader trend that

364 holds across the Triassic archosaur radiation. We posit that the qualitative NMDS ordination
365 method gives us the necessary lens for testing this hypothesis.

366

367 **Conclusions**

368 Simple quantitative measures of tooth shape were of limited use in characterizing the
369 Middle Triassic Manda Beds archosauriform tooth assemblage because of the highly conserved
370 morphology of many specimens. Instead, an ordination based on discrete characters provided a
371 more effective means of differentiating the teeth of distinct taxa. Nevertheless, we found little
372 evidence for significant ecological differentiation of tooth shape between the five taxa included
373 in our study. Most isolated teeth ($n = 17/21$) fall within the spectrum of recognized taxon
374 variation, and the remainder represent either unsampled taxa or unsampled intraspecific
375 variation.

376 Our relatively simple metrics can be used to describe subtle differences in tooth
377 morphology. These objective methods for grouping teeth provide a complimentary method for
378 assigning teeth to dietary roles, a practice that typically relies on qualitative comparisons to the
379 teeth of extant taxa of known diet (e.g. Fraser and Walkden, 1983; Sander, 1999; Barrett, 2000;
380 Hungerbühler, 2000) or other fossil taxa (e.g. Dzik, 2003; Hendrickx, Mateus & Araújo, 2015;
381 de Oliveira and Pinheiro, 2017). Furthermore, the methods applied herein provide an evaluation
382 of ecological disparity that is separate from the features used in phylogenetic analyses, so that we
383 can compare these two evolutionary phenomena independently. This method is readily
384 transferable to tooth assemblages from other localities pertaining to any vertebrate clade. Our
385 next step will be to apply this technique to richer Middle Triassic sites, as well as Late Triassic

386 sites, to understand how morphological and ecological diversity changed during the early stages
387 of the archosaur radiation.

388

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Figure 1

A sample of the *in situ* dental material used for baseline measurements in this study.

A - *Parringtonia* (NMT RB426) left dentary (left) in lateral (top) and medial (bottom) views and right dentary (right) in lateral and medial views. B - undescribed archosauriform taxon (NMT RB187) right maxilla in lateral and medial view. C - *Nundasuchus* (NMT RB48) holotype right dentary in lateral and medial views. D - *Asilisaurus* (NMT RB 837) (from left to right) right dentary in lateral and medial views, left maxilla in lateral and occlusal views, and right maxilla in lateral and occlusal views. Abbreviations: ap, ascending process of the maxilla; ds, dentary symphysis; ga, gastralnia; mg, Meckelian groove; mt III, metatarsal III; nf, nutrient foramen; pp, palatal process. All scale bars 1 cm.

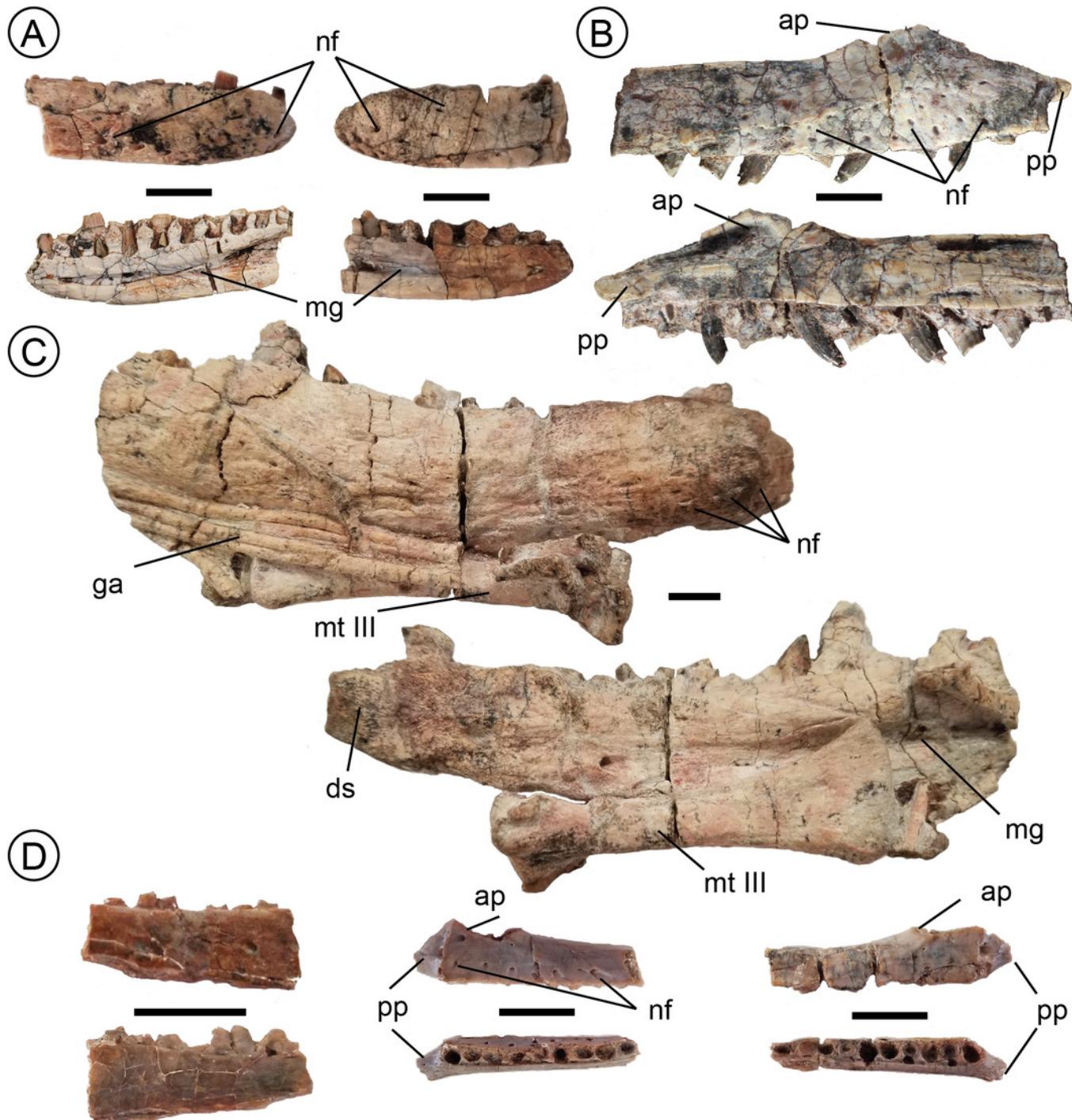


Figure 2

Examples of isolated teeth from the Manda Beds tooth assemblage.

A - Morphotype A specimens from left to right NMT RB807, NMT RB827, NMT RB809 in lateral and mesial views. Scale bars 1 cm. B - Morphotype B, specimens from left to right, NMT RB810, NMT RB819, NMT RB811 in lateral and mesial views. Scale bars 5 mm. C - sole representative of Morphotype C NMT RB831 in lateral and mesial views. Scale bar 1 cm. D - isolated teeth of known taxa. Top - *Nundasuchus* NMT RB48. Scale bar 1 cm. Bottom - *Parringtonia* NMT RB426 in lateral and mesial views. Scale bar 2 mm.

(A)



(B)



(C)



(D)



Figure 3

Visualization of discrete traits.

In all traits score 0 on left and score 1 on right. A - Trait 1, degree of recurvature, NMT RB819 (left) and NMT RB827 (right). B - Trait 2, fluting, NMT RB809 (left) and NMT RB426 (right). C - Trait 3, labiolingual curvature, NMT RB811 (left) and NMT RB819 (right). D - Trait 4, mesial margin angle, NMT RB811 (left) and NMT RB827 (right). E - Trait 5, labiolingual bulge, NMT RB811 (left) and NMT RB48 (right). F - Trait 6, dental caudae, NMT RB809 (left) and NMT RB810 (right). G - Trait 7, mesial vs distal serration series length, NMT RB831 (left) and NMT RB810 (right). H - Trait 8, denticle density per mm, NMT RB809 (left) and NMT RB810 (right), black lines equal 1 mm. I - Trait 9, mesial margin alignment, NMT RB810 (left) and NMT RB809 (right). J - Trait 10, mesial vs distal denticle density, NMT RB810 (left) and NMT RB809 (right). K - Trait 11, denticle shape variation along crown, NMT RB810 (left) and NMT RB48 (right).

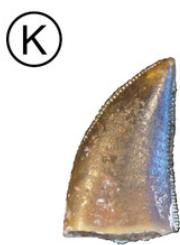
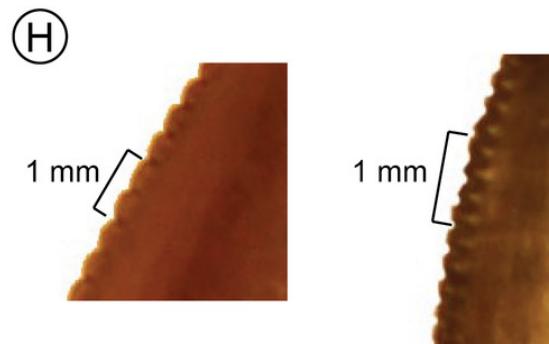
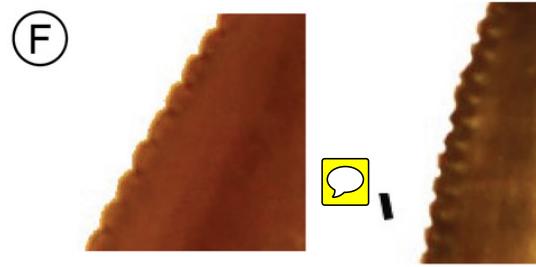
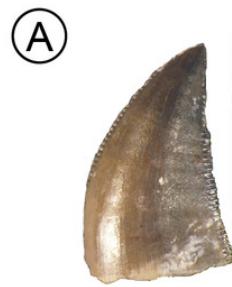


Figure 4(on next page)

Disparity of teeth measured by sum of variance

Disparity divided by taxon or morphotype. In the case of *Parringtonia* the two isolated teeth are an order of magnitude larger than the *in situ* teeth of this taxon, so this taxon was split. The sample sizes reflect the number of teeth with at least one of three measurements that was used to generate the predictive intervals.

Disparity

0.15

0.10

0.05

0.00

Asilisaurus

n = 13

NMT RB187

n = 13

Nundasuchus

n = 13

Pallisteria

n = 11

Parringtonia

n = 12

Parringtonia large

n = 2

Morphotype A

n = 14

Morphotype B

n = 5

Tooth Type

Figure 5(on next page)

Relationship between height and base shape of teeth divided by taxon.

The taxonomically unidentified teeth fall within a variety of the morphospaces generated by known taxa, rendering unambiguous referrals impossible. Some genera exhibit much greater variation in base shape ratio than others, potentially indicating a greater level of within-taxon variation.

Tooth Base Ratio

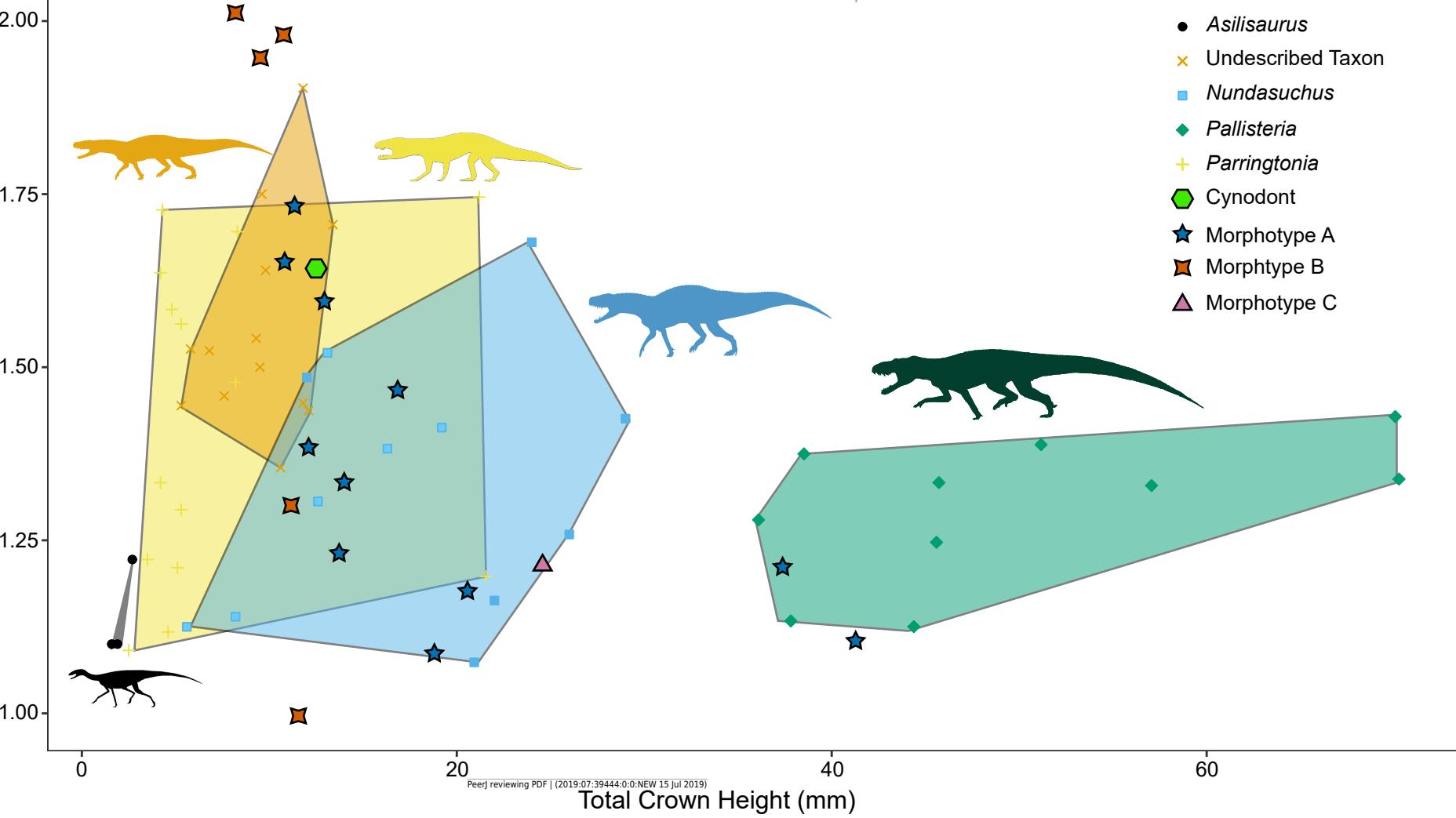


Figure 6(on next page)

Relationship between base width and fore-aft base length divided by taxon.

The overall ratio of base shape appears to be highly conserved with little deviation from the general trend. Differentiation between genera appears to be driven primarily by size.

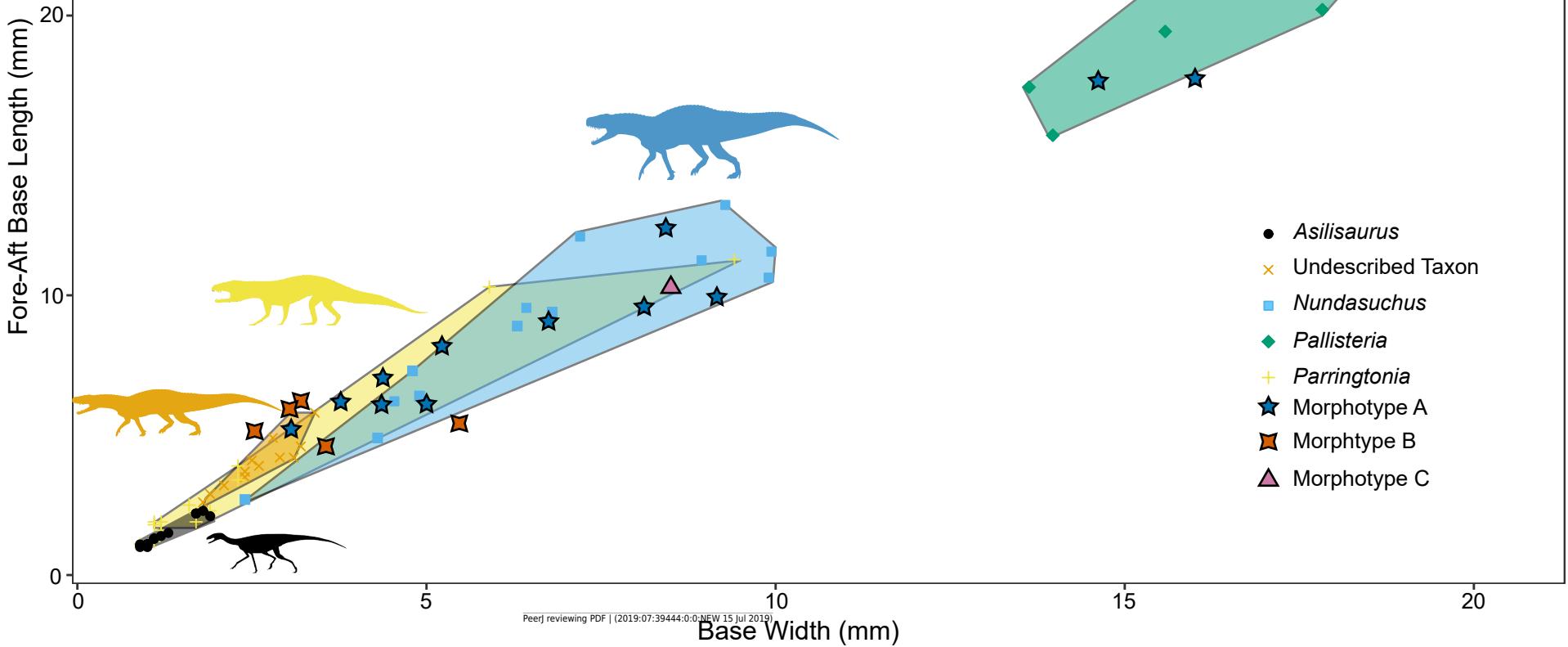


Figure 7(on next page)

Ordination plot of first two major NMDS axes of tooth morphospace.

Colored, transparent polygons represent the convex hulls of known taxa. Each point represents a separate tooth scoring. *Parringtonia* and NMT RB187 (undescribed taxon) share almost the same morphospace and there is substantial overlap between *Nundasuchus* and '*Pallisteria*' also. Morphotype A appears to be more variable than Morphotype B, which is clustered closer together within a subsection of overall Morphotype A morphospace. The proximity of *Asilisaurus* to *Nundasuchus* and '*Pallisteria*' is likely an artifact of incomplete scorings for *Asilisaurus* teeth.

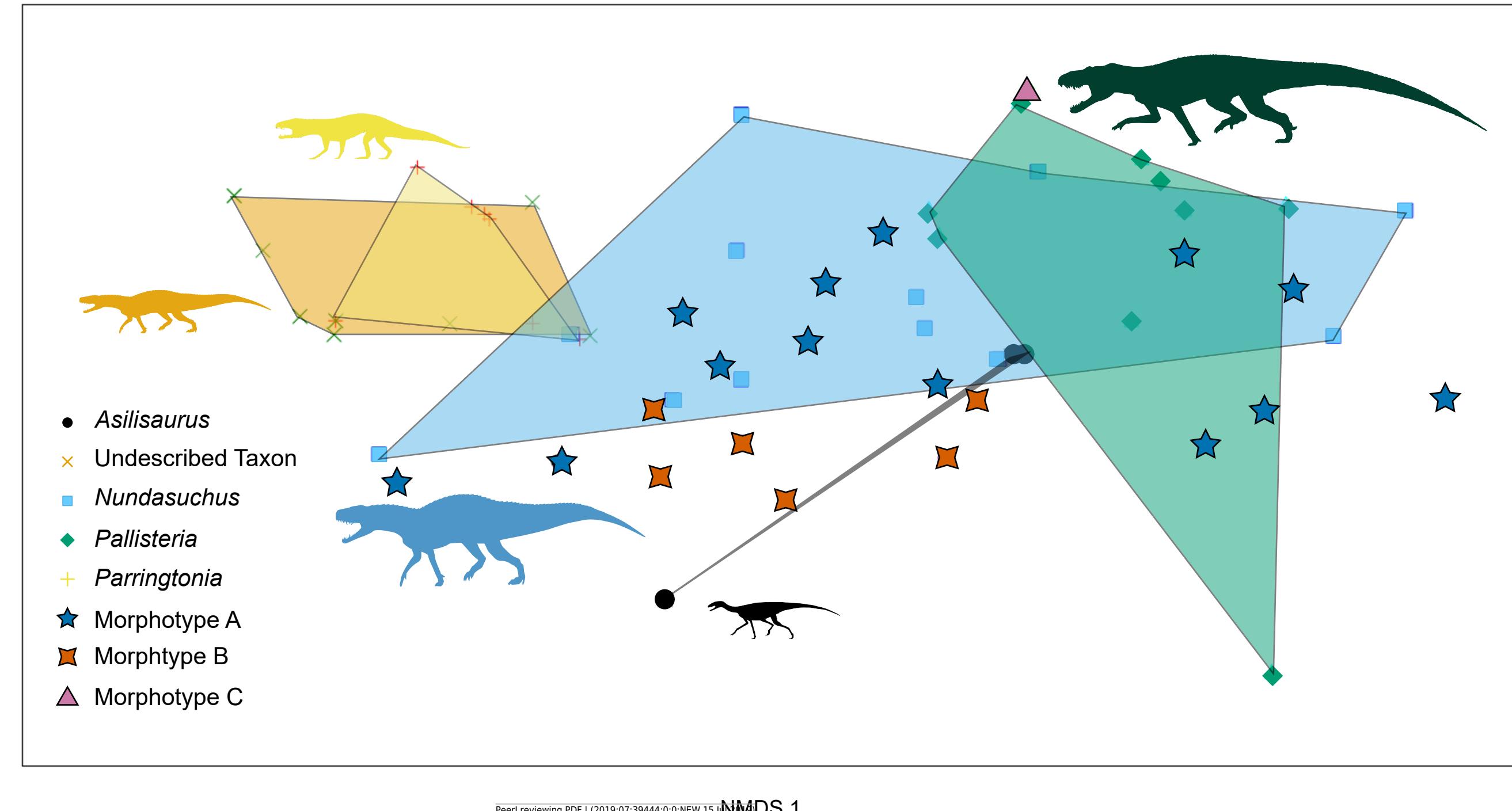


Figure 8(on next page)

Ordination plot of first two primary NMDS axes of tooth "averages" morphospace.

Taxa scoring represent "average" scores for each taxon. Only Morphotype C is represented by a single tooth. When using typical scores for taxa we find *Asilisaurus* is no longer near *Nundasuchus* and '*Pallisteria*' morphospace, but on the far side of ordination space.

- *Asilisaurus*
- ✖ Undescribed Taxon
- *Nundasuchus*
- ◆ *Pallisteria*
- ✚ *Parringtonia*
- ★ Morphotype A average
- ❖ Morphotype B average
- ▲ Morphotype C

NMDS 2

NMDS 1

Table 1(on next page)

Discrete character descriptions

Summary of the discrete, binary traits used for scoring teeth in the NMDS analysis.

| | Description |
|----|---|
| 1 | Tooth apex, location, relative to the distal margin of the tooth base: tip mesial to or in the same vertical plane as the distal edge (0) or tip is located more distal than the distal edge (=recurved) (1) |
| 2 | Tooth lingual/labial, surfaces: texture is smooth (lack of crenulations, ridges, etc.) (0) or surface texture possess a series of parallel ridges from tooth apex to base (=fluted) (1) |
| 3 | Tooth labial/lingual, shape: crown curvature unequal (one side expanded relative to other) (0) or equal labial and lingual curvature (1) |
| 4 | Mesial tooth margin, shape: curvature angles change gradually (0) or angle changes abruptly at a single discrete point along mesial edge (1) |
| 5 | Tooth crown, size: labiolingual widths dorsal to the tooth crown base are all less than the crown base width (0) or a crown labiolingual width dorsal to the tooth crown base is greater than the crown base width (1) |
| 6 | Mesial/distal crown margins, surfaces: denticle caudae (= grooves on crown surface from between individual denticles) are absent (0) or present (1) (from Abler, 1992) |
| 7 | Mesial margin, length: mesial denticle row ends at a point sub-equal with distal denticle row (0) or mesial denticle row ends significantly further apically on crown than distal row (1). Can only be scored for teeth with both mesial and distal denticle series. |
| 8 | Mesial/distal margins, denticle density: number of mesial and distal denticles is < 3 per mm (0), or greater than or equal to 3 per mm (1). Measurements are taken near the middle of the carina. |
| 9 | Mesial margin, location: vertical axis of the mesial carina is in line the mesial-distal long axis (0) or laterally offset from the mesial distal long axis (1) |
| 10 | Mesial/distal margins, size: average size of mesial and distal denticles are the same (0) or the average size of the mesial and distal denticles is different (1) |
| 11 | Mesial/distal margins, shape: lateral profile shape of mesial and distal denticles remains constant (0) or denticles' lateral profile changes shape (e.g. rounded to square) (1) |

Table 2(on next page)

Results of linear model (base ~ total crown height + taxon)

All measures of significance are calculated in reference to the intercept, *Asilisaurus*.

Therefore, while the undescribed pseudosuchian and *Parringtonia* can be differentiated in the model from *Asilisaurus*, the interrelationships are unknown.

| | Estimate | Standard Error | t-value | p-value |
|--------------------------------|----------|----------------|---------|---------|
| <i>Asilisaurus</i> (intercept) | 1.1285 | 0.1011 | 11.156 | <0.0001 |
| Total Crown Height (mm) | 0.0059 | 0.0035 | 1.714 | 0.0933 |
| Undescribed | 0.3718 | 0.1148 | 3.237 | 0.0022 |
| <i>Nundasuchus</i> | 0.0995 | 0.1247 | 0.798 | 0.4290 |
| ' <i>Pallisteria</i> ' | -0.1249 | 0.2007 | -0.622 | 0.5369 |
| <i>Parringtonia</i> | 0.2490 | 0.1127 | 2.210 | 0.0321 |

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Table 3(on next page)

Pairwise comparisons of taxa used in the linear model.

The undescribed pseudosuchian is readily differentiable from most taxa, with the exception of *Parringtonia*. Confidence intervals were generated using a 95% confidence level.

| Taxon | lsmeans | Standard Error | df | Lower CL | Upper CL |
|------------------------|---------|----------------|----|----------|----------|
| <i>Asilisaurus</i> | 1.2354 | 0.1150 | 46 | 1.0039 | 1.4669 |
| Undescribed | 1.6071 | 0.0568 | 46 | 1.4929 | 1.7214 |
| <i>Nundasuchus</i> | 1.3348 | 0.0505 | 46 | 1.2332 | 1.4365 |
| ' <i>Pallisteria</i> ' | 1.1105 | 0.1225 | 46 | 0.8640 | 1.3570 |
| <i>Parringtonia</i> | 1.4844 | 0.0595 | 46 | 1.3645 | 1.6042 |

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| Contrast | Estimate | SE | df | t ratio | p-value |
|--|----------|--------|----|---------|---------|
| <i>Asilisaurus</i> – Undescribed | -0.3718 | 0.1148 | 46 | -3.237 | 0.0181 |
| <i>Asilisaurus</i> – <i>Nundasuchus</i> | -0.0995 | 0.1247 | 46 | -0.798 | 0.9299 |
| <i>Asilisaurus</i> – 'Pallisteria' | 0.1249 | 0.2007 | 46 | 0.622 | 0.9708 |
| <i>Asilisaurus</i> – <i>Parringtonia</i> | -0.2490 | 0.1127 | 46 | -2.210 | 0.1944 |
| Undescribed – <i>Nundasuchus</i> | 0.2723 | 0.0751 | 46 | 3.625 | 0.0062 |
| Undescribed – 'Pallisteria' | 0.4967 | 0.1571 | 46 | 3.162 | 0.0222 |
| Undescribed – <i>Parringtonia</i> | 0.1228 | 0.0677 | 46 | 1.813 | 0.3788 |
| <i>Nundasuchus</i> – 'Pallisteria' | 0.2244 | 0.1343 | 46 | 1.671 | 0.4614 |
| <i>Nundasuchus</i> – <i>Parringtonia</i> | -0.1495 | 0.0770 | 46 | -1.942 | 0.3107 |
| 'Pallisteria' – <i>Parringtonia</i> | -0.3739 | 0.1631 | 46 | -2.292 | 0.1659 |

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