

# Deep-time biodiversity patterns and the dinosaurian fossil record of the Late Cretaceous Western Interior, North America

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1 Deep-time Biodiversity Patterns and the Dinosaurian Fossil Record of the Late  
2 Cretaceous Western Interior, North America

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9 **Abstract**

10 In order for palaeontological data to be informative to ecologists seeking to understand the causes  
11 of today's diversity patterns, palaeontologists must demonstrate that actual biodiversity patterns  
12 are preserved in our reconstructions of past ecosystems. During the Late Cretaceous, North America  
13 was divided into two landmasses, Laramidia and Appalachia. Previous work has suggested strong  
14 faunal provinciality on Laramidia at this time, but these arguments are almost entirely qualitative.  
15 We quantitatively investigated faunal provinciality in ceratopsid and hadrosaurid dinosaurs using a  
16 biogeographic network approach and investigated sampling biases by examining correlations  
17 between dinosaur occurrences and collections. We carried out a model-fitting approach using  
18 generalized least-squares regression to investigate the sources of sampling bias we identified. We  
19 find that while the raw data strongly support faunal provinciality, this result is driven by sampling  
20 bias. The data quality of ceratopsids and hadrosaurids is currently too poor to enable fair tests of  
21 provincialism, even in this intensively sampled region, which probably represents the best-known  
22 Late Cretaceous terrestrial ecosystem on Earth. To accurately reconstruct biodiversity patterns in  
23 deep time, future work should focus on smaller scale, higher resolution case studies in which the  
24 effects of sampling bias can be better controlled.

25

26 **Keywords:** faunal provincialism, endemism, Laramidia, sampling bias, Ceratopsidae, Hadrosauridae.

## 27 **Introduction**

28           In order to predict how biodiversity patterns on today's Earth will respond to climate  
29 change, the factors that cause biodiversity distributions must be understood [1,2]. Deep-time  
30 perspectives can provide novel insights into the controls on biodiversity distribution. By examining  
31 biodiversity distributions at times in Earth's history when climate, continental arrangement, and  
32 oceanic currents were different than today, ecological hypotheses about the causative mechanisms  
33 behind biodiversity distribution and the establishment of modern patterns can be tested [3–7].  
34 However, if palaeontologists wish their data to be informative to those working on the causative  
35 mechanisms of modern-day biodiversity patterns, we must first demonstrate that actual biodiversity  
36 patterns are preserved in our reconstructions of past ecosystems, and that we are able to overcome  
37 the many sampling biases that affect the fossil record [e.g., 8–10].

38           Dinosaurs are an exceptional model system for studying biodiversity and macroevolution in  
39 terrestrial vertebrates. For >150 million years, from the Late Triassic to the end of the Cretaceous,  
40 they dominated terrestrial ecosystems, occupied every continent, and radiated into a wide variety of  
41 ecological niches. Because of public interest, they are the best-sampled Mesozoic terrestrial  
42 vertebrate group, and their fossils have been collected for well over 150 years [11–14]. Arguably, the  
43 best-sampled part of the dinosaurian fossil record is the Late Cretaceous of the Western Interior  
44 region of North America [15–17]. During the Late Cretaceous, North America was divided into two  
45 landmasses, Laramidia to the west and Appalachia to the east, by the epicontinental Western  
46 Interior Seaway. In the latter stages of the Late Cretaceous (Campanian and Maastrichtian), large-  
47 bodied herbivorous niches in Laramidia were dominated by two groups of ornithischian dinosaurs,  
48 the hadrosaurs and ceratopsids. The body fossil record of the latter is entirely restricted to North  
49 America at this time, with the exception of a single taxon [18,19]. Study of these herbivorous  
50 dinosaurs has provided major insight into dinosaur behaviour, palaeoecology and biogeographic  
51 patterns [e.g., 20–32].

52 Numerous workers have argued for strong faunal provinciality in Laramidia throughout the  
53 Campanian, and have divided the landmass into northern and southern faunal provinces [e.g., 33–  
54 35]. This signal is particularly clear in chasmosaurine ceratopsids, where virtually all species are  
55 recognised from either northern or southern Laramidia, but not both [34–36]. This endemism in  
56 ceratopsids is thought to have driven high levels of diversity, underpinning their radiation [34]. Since  
57 no geological or geographical barrier has thus far been identified between northern and southern  
58 Laramidia [37], the boundary between the northern and southern provinces has been suggested to  
59 be related to latitudinal climate, temperature, or rainfall patterns [34, 35, 38] or was maintained due  
60 to competition between local populations [39]. The patterns of apparent provincialism decrease in  
61 the Maastrichtian, coincident with overall regression of the Western Interior Seaway [15, 17, 33, 36].

62 These hypotheses of biogeographic provincialism, however, remain controversial. With very  
63 few exceptions [35, 36], studies that advocate for provincialism are based on qualitative  
64 observations [e.g., 33, 34] and arise from comparisons of the fauna of specific geological formations  
65 [e.g., 33–35]. Recent research has, however, suggested that some of the formations used in such  
66 studies are not contemporaneous [17, 40] and that the length of time-intervals used results in the  
67 amalgamation of multiple successive faunas [17, 37]. Many studies advocating faunal endemism are  
68 based on taxonomic decisions that have proven controversial and the conclusions have been called  
69 into question as a result [e.g., 37, 39]. Additionally, it remains a possibility that faunal provinces  
70 within the Campanian are an artefact of sampling: most Campanian dinosaur occurrences are known  
71 from Alberta, Montana, southern Utah and northern New Mexico, with far less sampling having  
72 occurred in northern Utah and Wyoming [16].

73 Three quantitative studies have investigated the provincialism hypothesis in dinosaurs of the  
74 Late Cretaceous Western Interior. Gates *et al.* [35] used a variety of statistical techniques to assess  
75 similarity between Campanian northern and southern faunas, and found evidence for either two  
76 distinct provinces with a broad area of overlap between them, or a latitudinal diversity gradient. The

77 statistical techniques employed were unable to distinguish between these two hypotheses, and their  
78 results regarding dinosaurs were inconclusive. They suggested further investigation into the causes  
79 of dinosaur distribution in the Western Interior. Berry [36] used a phylogenetic approach to assess  
80 biogeography within the Campanian and found no evidence for endemic sub-clades of  
81 chasmosaurine ceratopsids, arguing that this would be expected if there was a major barrier to  
82 dispersal, or niche conservatism related to climate. Vavrek & Larsson [15] investigated faunal  
83 endemism in the Maastrichtian of Laramidia using measures of beta-diversity. After correcting for  
84 sampling biases they found little evidence of provincialism, instead suggesting a homogeneous  
85 dinosaurian fauna across the Western Interior region at the very end of the Cretaceous; however,  
86 they did not test to see whether apparent biogeographic patterns within the Campanian were also  
87 caused by sampling.

88         Herein, we quantitatively test hypotheses of faunal endemism in both the Campanian and  
89 Maastrichtian using biogeographic and multivariate statistical approaches. We focus our study on  
90 ceratopsid and hadrosaurian dinosaurs, as these megaherbivores have well understood phylogenies  
91 and taxonomies and have been at the centre of previous discussions of faunal provinciality in this  
92 region. The distinctiveness of northern and southern Laramidian provinces are tested using  
93 phylogenetically-corrected Biogeographic Connectedness (pBC). This quantitative method uses a  
94 network approach to assess phylogenetic distances between taxa in different geographic areas,  
95 resulting in a metric that quantifies the degree of faunal provinciality versus cosmopolitanism. It has  
96 been used successfully to understand changes in faunal compositions through the Carboniferous–  
97 Permian transition [41], and the Permian–Triassic and Triassic–Jurassic extinction events [42]. We  
98 also introduce additions to the methodology that address concerns regarding variation in sampling  
99 through time. To investigate the impact of sampling bias on our results, we examine correlations  
100 between occurrences (records of specimens) and collections (sites where specimens have been  
101 collected) with latitude, and use multivariate regression to examine which sources of sampling bias

102 best explain sampling patterns. We use our results to determine whether it is possible to identify  
103 true geographic patterns of biodiversity on a continental scale in this very well-sampled area.

104

## 105 **Methods**

106 *Taxon sampling & phylogeny*—Since no complete phylogenetic analysis of all ceratopsids is  
107 available, we built an informal supertree of all ceratopsid taxa considered valid in recent  
108 phylogenetic analyses by combining the phylogenetic results of [18] for chasmosaurines and [19] for  
109 centrosaurines. We resolved polytomies by removing *Nedoceratops*, a taxon some workers consider  
110 to be invalid [43, but see 44], from the data matrix in [18] and re-analysing the dataset. This resolved  
111 polytomies in the clade containing the common ancestor of *Eotriceratops*, *Triceratops* and all of its  
112 descendants. The resulting supertree includes 67 taxa and represents a consensus of current views  
113 on ceratopsian phylogeny (Fig. S1A).

114 The structure of the hadrosaurid tree is based on several key recent analyses [45–47]. We  
115 resolved polytomies and added taxa considered valid but not included in those references using  
116 other recent phylogenetic analyses [48–51]. The resulting supertree includes 55 taxa and represents  
117 a current reasonable estimate of hadrosaur phylogeny (Fig. S1B).

118 *Stratigraphic age and geographic data*—Age for North American hadrosaur and ceratopsid  
119 species was obtained from the primary literature. The formations in which taxa occurred were found  
120 from the Paleobiology Database (PBDB; [www.paleobiodb.org](http://www.paleobiodb.org)), and the most recent absolute age  
121 estimate of those formations was obtained from the primary literature (see Online Supplementary  
122 Material, OSM, for sources). The age and geographic data for taxa outside of North America was  
123 obtained from the Paleobiology Database. pBC requires a *priori* assignment of geographic regions to  
124 test hypotheses of biogeographic connectedness, so we assigned dinosaurs to either northern  
125 Laramidia or southern Laramidia. Northern Laramidia includes taxa found in Wyoming and further

126 north; southern Laramidia includes taxa found in Utah and further south, following previous studies.  
127 Age data was used to time-calibrate the phylogenetic trees using the ‘timePaleoPhy’ function of the  
128 Strap package [52] in R version 3.5.2 [53] with the minimum branch length option specified  
129 (type=‘mbl’). Whilst it would be ideal to use high-resolution bins to test patterns of biogeography  
130 through the Late Cretaceous [17], too few taxa would be present in each bin to permit the use of  
131 pBC. Consequently, we divided taxa into Campanian and Maastrichtian time bins, which also has the  
132 benefit of allowing for comparison between previous studies of faunal provincialism in this area.  
133 Where a taxon’s stratigraphic range/uncertainty crossed the Campanian–Maastrichtian boundary, it  
134 was included in both time intervals.

135 *pBC*—We calculated pBC for Campanian ceratopsids, Maastrichtian ceratopsids, and  
136 Campanian hadrosaurs. Sampling of Maastrichtian hadrosaurs was too sparse, particularly in  
137 southern Laramidia, to calculate meaningful pBC values. Trees were pruned to exclude taxa from  
138 timeslices other than the one being analysed, and were made ultrametric prior to analysis. pBC was  
139 calculated using the function BC of the package ‘dispeRse’ (available at  
140 [github.com/laurasoul/disperse](https://github.com/laurasoul/disperse)). We initially varied the constant  $\mu$  [see 41] from 1–15 million years;  
141 subsequent analyses used a constant  $\mu$  of 10. Data were jack-knifed 1000 times to produce a  
142 distribution of possible pBC values. To address concerns about the potential for a relationship  
143 between pBC and taxon sample size [54], we calculated rarefaction curves for pBC for the  
144 ceratopsian data (to facilitate comparisons between the Campanian and Maastrichtian). Sample  
145 sizes were rarefied down to a minimum number of five taxa. 95% confidence intervals for the  
146 rarefaction curves were generated using 1000 replicates at each sampling level.

147 *Randomization of data (null model)*—In order to determine whether pBC for each clade and  
148 time interval was significantly different from random, we randomly permuted the geographic areas  
149 in which taxa are found. We generated 1000 permutations of the data for each clade and time

150 interval and calculated pBC for each permutation. The pBC for the unpermuted data was compared  
151 to the distribution of permuted pBC values to establish statistical significance ( $p < 0.05$ ).

152         *Sampling bias*—To investigate whether biogeographic patterns we observed in the pBC  
153 results were influenced by sampling bias, we downloaded raw occurrence data for ceratopsids and  
154 hadrosaurs for the Campanian and Maastrichtian from the PBDB. We then downloaded North  
155 American dinosaur-bearing collections and North American tetrapod-bearing collections for each  
156 timeslice, and plotted occurrences and collections with latitude. We compared the curves using  
157 Spearman's rho and Kendall's tau.

158         To investigate the possible causes of sampling bias we identified, we statistically examined  
159 correlations between occurrences and outcrop area, depositional environment, and proxies for  
160 exposure. First, we imported publicly available USGS state-level and Canadian Province digital  
161 geological maps ([www.ngmhd.usgs.gov](http://www.ngmhd.usgs.gov); <https://ags.aer.ca/publication/map-600>;  
162 <https://geohub.saskatchewan.ca/datasets/bedrock-geology>) into ArcMap 10 ([www.esri.com](http://www.esri.com)),  
163 identified Campanian and Maastrichtian strata, and assigned an environmental attribute  
164 determining whether strata were deposited in a terrestrial, marine, or mixed setting (OSM). These  
165 data, along with maximum green vegetation fraction (MGVF) and slope, both proxies for exposure,  
166 were imported into R (version 3.5.0). Methods for generating MGVF and slope are provided in OSM.  
167 Level plots of total outcrop area, terrestrial, mixed and marine outcrop area, slope, and MGVF were  
168 produced using the `levelplot()` function of the `rasterVis()` package [55] (Fig. S2).

169         To investigate the power of each or a combination of these variables to explain the dinosaur  
170 occurrence data, we carried out a model-fitting approach using generalized least-squares regression  
171 (GLS). Ceratopsian and hadrosaur occurrences from the Paleobiology Database were counted in each  
172 1-degree latitudinal bin (latitude is modern latitude, rather than palaeolatitude). Models compared  
173 latitudinal changes in ceratopsian and hadrosaur occurrences to changes in four different measures  
174 of outcrop area (see OSM), MGVF and slope. GLS autoregressive models were fitted to combinations

175 of potential explanatory variables. We used with a first order autoregressive model (corARMA) fitted  
176 to the data to account for spatial autocorrelation using the function `gls()` in the R package `nlme` v.  
177 3.1–150 [56]. GLS reduces the chance of overestimating statistical significance of regression lines  
178 due to serial correlation in the latitudinal series.

179 Data series were ln-transformed prior to analysis to ensure normality and homoskedasticity  
180 of residuals. We calculated likelihood-ratio based pseudo- $R^2$  values using the function `r.squaredLR()`  
181 of the R package `MuMIn` [57]. Results were compared using Akaike's information criterion for small  
182 sample sizes (AICc) and Akaike weights were calculated to identify the best combination of  
183 explanatory variables from those tested. AICc was calculated using the function `AICc()` of the R  
184 package `qpcR` [58], and Akaike weights calculated using the `aic.w()` function of the R package  
185 `phytools` [59].

#### 186 *Sampling bias and pBCs*

187 To test the impact of the Campanian bimodal sampling distribution on pBC results, we ran a  
188 second pBC test where we randomly removed 95% of ceratopsian taxa from the Maastrichtian that  
189 occurred between 35 and 50 degrees of latitude. We chose these latitudinal boundaries to enforce a  
190 similar bimodal latitudinal diversity gradient on the Maastrichtian data as seen in the Campanian  
191 (see Results). The remaining distribution of occurrences was used to re-run pBC analyses (with a  $\mu$  of  
192 10), and this process was repeated 1000 times for increased accuracy of results. pBC scores were  
193 recorded for each run, and the resulting distribution was used to calculate the mean pBC to compare  
194 against the original Maastrichtian ceratopsian pBC score and produce a probability density curve to  
195 estimate the probability of different values of pBC scores.

196

## 197 **Results**

198           The observed value of pBC for Campanian ceratopsids was 0.05, while that for Campanian  
199 hadrosaurs was 0.11, and for Maastrichtian ceratopsids the observed value was 0.16. pBC was  
200 therefore lower for ceratopsids in the Campanian than in the Maastrichtian, and endemism was  
201 correspondingly higher, in agreement with previous studies [33, 36]. Jack-knifed distributions of  
202 ceratopsid pBC for the Campanian and Maastrichtian overlap (Fig. 1A–C), but their median values  
203 are strongly significantly different from each other (Wilcox Test,  $W=60235$ ,  $p=0.00$ ). Rarefaction  
204 curves for ceratopsids for the Campanian and Maastrichtian indicate a much higher pBC in the  
205 Maastrichtian than in the Campanian at equivalent levels of sampling, although the confidence  
206 intervals do overlap, particularly at lower sampling levels (Fig. 1D). This demonstrates that the higher  
207 pBC of the Maastrichtian is not a consequence of sampling lower numbers of species in that interval  
208 in comparison to the Campanian. Higher pBC equates to more cosmopolitan faunas, and thus this  
209 result supports lower endemism in Laramidia during the Maastrichtian when compared to the  
210 Campanian.

211           Values for pBC for both Campanian and Maastrichtian ceratopsid data are statistically  
212 significantly lower than for datasets in which the geographic areas are randomized (Campanian,  
213  $p=0.00$ ; Maastrichtian,  $p=0.015$ ; Fig. S3), and the same is true for the Campanian hadrosaur data  
214 ( $p=0.00$ ; Fig. S3). This indicates that endemism was statistically significantly higher than in all  
215 randomized datasets across both time intervals, and supports previous qualitative hypotheses of  
216 distinct northern and southern provinces in Laramidia [e.g., 33–35]

217           Curves of raw occurrence data with latitude for hadrosaurs and ceratopsids in both the  
218 Campanian and the Maastrichtian correlate strongly and statistically significantly with both dinosaur-  
219 bearing and tetrapod-bearing collections (Fig. 2; Fig. S4; OSM). During the Campanian, sampling and  
220 occurrences are focused at two latitudes: 51-49 degrees north, which corresponds with the Dinosaur  
221 Park, Oldman and, to a lesser extent, the Foremost formations, and 36-37 degrees north, which  
222 corresponds primarily with the Kirtland/Fruitland, Aguja and Kaiparowits formations (Fig. 2A, B; Fig.

223 S4A, B). These two areas have been sampled orders of magnitude better than the surrounding  
224 latitudinal bins [16], although there are tetrapod- and dinosaur-bearing formations across the  
225 majority of the Western Interior at this time (Fig. 2A, B; Fig. S4A, B). In the Maastrichtian, sampling is  
226 more evenly spread across the range of latitudes for which we have hadrosaur and ceratopsid body  
227 fossils (Fig. 2C, D; Fig. S4; C, D; [16]). These data are strongly indicative that the provinciality  
228 observed based on raw data in the Campanian could be due to intensive sampling in the Dinosaur  
229 Park Formation and Kirtland/Fruitland Formations with a lack of sufficient sampling between, and  
230 our observed increase in pBC (=reduced endemism) in the Maastrichtian is due to increased  
231 latitudinal coverage of sampling.

232 The mean pBC score of Maastrichtian ceratopsians subjected to a Campanian-style sampling  
233 distribution was 0.0351 with a standard deviation of 0.0476, significantly lower than the original pBC  
234 score of 0.16. The probability of a pBC score less than or equal to 0.8 was 0.78 (OSM and Fig. S5).  
235 These results provide a further indication that sampling bias is driving pBC scores of Campanian  
236 fauna

237 A lack of sampling in the area between 49 degrees north and 37 degrees north (the  
238 'sampling peaks') in Campanian strata could be caused by a variety of factors. It has long been  
239 known that rock outcrop area is strongly correlated with raw diversity [e.g., 60, 61,]; if there is less  
240 outcrop, there are fewer opportunities for palaeontologists to sample the rocks, and fewer fossils  
241 found as a consequence. As terrestrial organisms, the vast majority of dinosaur fossils are found in  
242 formations that were deposited on land. If Campanian rocks between the sampling peaks are  
243 primarily marine, there will be fewer opportunities for dinosaur fossils to be preserved, and thus  
244 fewer opportunities for them to be sampled by palaeontologists. Fossils are primarily found where  
245 bare rock is exposed at the surface. If less rock is exposed between the sampling peaks than in the  
246 areas of the peaks themselves, there will be fewer opportunities for fossils to come to light.

247           GLS analyses recovered the following best models (highest AICc weights) for outcrop and  
248 tetrapod occurrence masks (see OSM for additional results): Campanian hadrosaurs, summed  
249 outcrop area + MGVF + slope; Campanian ceratopsians, non-marine total outcrop area;  
250 Maastrichtian hadrosaurs, null model; Maastrichtian ceratopsians, null model. However, in nearly all  
251 cases the correlations are non-significant (OSM) and only the Campanian hadrosaur model results  
252 had a strong overall explanatory power (OSM). This indicates that the potential sampling bias with  
253 latitude in the Campanian that we have identified cannot be fully explained by any of these variables  
254 and other sources of sampling bias that are hard to quantify may additionally be responsible.

255

## 256 **Discussion**

257           Several authors have suggested that the apparent faunal provincialism in Laramidia during the  
258 Late Cretaceous is an artefact, either because the formations in which dinosaurian taxa have been  
259 found are not contemporaneous [17, 37, 40] or due to uneven sampling of the fossil record [15, 16].  
260 Our results show that while the raw data clearly supports faunal endemism, particularly in the  
261 Campanian, this pattern is driven by a lack of sampling outside of two specific latitudinal belts on  
262 Laramidia (51-49 degrees north, which corresponds with the Dinosaur Park, Oldman and Foremost  
263 formations, and 36-37 degrees north, which corresponds primarily with the Kirtland/Fruitland,  
264 Aguja and Kaiparowits formations). This sampling bias cannot be fully explained by differences in  
265 outcrop area across the region, or by differences in slope or vegetation, which are factors that affect  
266 rock exposure.

267           There are numerous other factors that can bias sampling, but these are very difficult to quantify.  
268 Low sampling between the northern and southern sampling peaks could occur if palaeontologists  
269 have yet to prospect the area to the same degree that they have in the north and south. The Late  
270 Cretaceous of the Western Interior has been intensively sampled for dinosaur fossils for over 100

271 years, and it is now probably the best-known Late Cretaceous ecosystem anywhere on Earth [15,16].  
272 It is therefore highly unlikely that large parts of it remain unexplored for dinosaurs, and the fact that  
273 dinosaur fossils are known from the area between the sampling peaks during the Maastrichtian  
274 suggest the area has been explored. Lack of exploration for fossils is therefore unlikely to be the  
275 primary driver of the uneven sampling patterns we have observed.

276 The 'common cause hypothesis' [e.g., 62] suggests that correlations between raw diversity and  
277 sampling proxies (e.g., numbers of formations) are driven by a third factor, usually sea level.  
278 Although initially formulated for marine environments, the possibility of a sea-level driven common  
279 cause on land has also been discussed [e.g., 63]. During sea-level high stands sediment flux to inner  
280 shelves and marginal marine areas is high; this results in both high potential for the preservation of  
281 fossils due to rapid burial and high diversity due to habitat fragmentation leading to endemism and  
282 increased beta diversity. Conversely, sediment bypasses inner shelf environments during low stands,  
283 reducing sediment flux and leading to poorer preservation of fossils due to a lower chance of rapid  
284 burial, while diversity is lower due to cosmopolitanism. Although the effect of eustatic sea level  
285 changes on the global terrestrial fossil record of vertebrates has been questioned [63], Chiarenza et  
286 al. [16] demonstrated that the areas of our northern and southern sampling peaks correlated with  
287 high sediment fluxes and low runoff rates during the Campanian. It is therefore possible that  
288 reduced sampling between our sampling peaks is because this area was less suitable for fossil  
289 preservation in the Campanian. Indeed, Chiarenza *et al.* [16] suggested that faunal provincialism in  
290 the Campanian was a sampling bias at least partially due to variation in climatic induced taphonomic  
291 suitability between northern and southern regions.

292 Historical collecting practises and/or land ownership might also play a role in the sampling  
293 patterns we have observed. If the proportion of outcrop on public land was reduced in the areas  
294 outside of our sampling peaks, this might mean palaeontologists have less access to explore there  
295 for fossils. Furthermore, if there is a particularly field-active palaeontological institution close to an

296 area of Campanian outcrop, or long-term agreements in place with landowners, this may have  
297 allowed prospecting to occur more regularly over a longer period of time in specific areas. A bias  
298 may also be introduced by uneven regional entry of data into online databases such as the PBDB.  
299 Such a bias could stem from monographs on specific formations or museums whose collections  
300 focus on specific areas that also have online databases. Data from these sources are comparatively  
301 easy to enter into the PBDB and thus could be contributing to the sampling patterns we observed.

302 It seems highly likely that a combination of available outcrop area, rocks suitable for the  
303 fossilization of vertebrate remains, and an interplay between climate, topography and historical  
304 collection and data entry practises is responsible for variations in sampling across the Western  
305 Interior, which have resulted in apparent northern and southern faunal provinces on Laramidia.

#### 306 *Taxonomic differences in northern and southern Laramidia*

307 Despite the fact that we find faunal provincialism in the Late Cretaceous Western Interior to  
308 mostly likely be due to sampling bias based on currently available data, it is clear that different taxa  
309 are found in the northern and southern areas of Laramidia [33–35]. This is especially clear in  
310 chasmosaurine ceratopsids, where there is almost no overlap at all between taxa found in the north  
311 and those found in the south [34, but see 36, 39]). It has been demonstrated that many of these taxa  
312 were not contemporaneous [40], which would at least partially explain taxonomic differences. But,  
313 in addition, the study area covers 12 degrees of latitude and climate would have varied significantly  
314 over that area, even in a greenhouse world where latitudinal temperature gradients were reduced  
315 relative to today [39, 64]. General circulation models for the Campanian show significant variation in  
316 mean annual temperature and rainfall patterns with latitude across Laramidia [16] and recent  
317 research has suggested elevated temperature gradients in a climatic transition zone between the  
318 northern and southern faunal provinces [38]. Given that there is evidence for both spatial [26] and  
319 functional [29] niche partitioning in Laramidia's large herbivores, taxonomic differences between the  
320 north and south could be related to climatic preference, and there may well have been a latitudinal

321 biodiversity gradient across the area. Unfortunately, we have demonstrated here that that raw data  
322 is currently too influenced by sampling biases for such biodiversity patterns to be reconstructed.

323

## 324 **Conclusion**

325 We show that data quality of Campanian and Maastrichtian ceratopsids and hadrosaurs, two  
326 of the most abundant clades of dinosaurs in the Late Cretaceous of North America, is currently too  
327 poor to enable fair tests of endemism and provincialism. In order to effectively test hypotheses  
328 regarding the causative mechanisms of biodiversity distribution, palaeontologists must demonstrate  
329 either that the fossil record preserves true biodiversity patterns at high levels of temporal  
330 resolution, or that methods exist that can adequately overcome sampling biases. The Western  
331 Interior region represents probably the most densely-sampled Late Cretaceous terrestrial region  
332 worldwide [15, 16], but even in this intensively-sampled area, it is not currently possible to  
333 reconstruct diversity patterns at the regional scale. In order for palaeontologists to make a  
334 meaningful contribution to ecological hypotheses about future biodiversity change, we must focus  
335 our efforts on smaller-scale case studies, where temporal resolution is high, stratigraphic correlation  
336 is well-established, and where sampling biases are likely to be more homogenous and can be more  
337 easily quantified. A good example of a recent such study is [65]. The results of multiple high-  
338 resolution case studies can then be compared globally to establish the rules that governed past  
339 biodiversity distributions.

340

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347

#### 348 **Author contributions**

349 SCRM & RJB designed the study. SCRM, RJB & CDD wrote the manuscript. RIM & SCRM collected  
350 data and generated supertrees. SCRM, RJB and CDD ran analyses. All authors provided critical  
351 comments on the manuscript.

352

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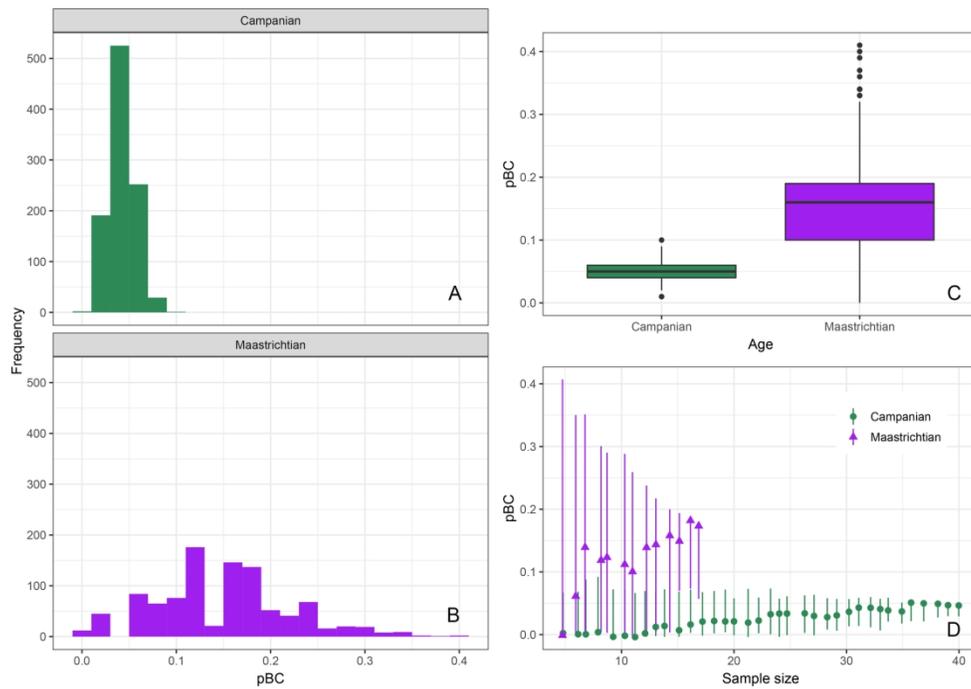


Figure 1. A–C. Jackknifed distributions of pBC values for Campanian (green) and Maastrichtian (purple) ceratopsids. D. Rarefaction curves for Campanian (green circles) and Maastrichtian (purple triangles) ceratopsids. Error bars show the 95% confidence intervals of values obtained during rarefaction.

175x123mm (300 x 300 DPI)

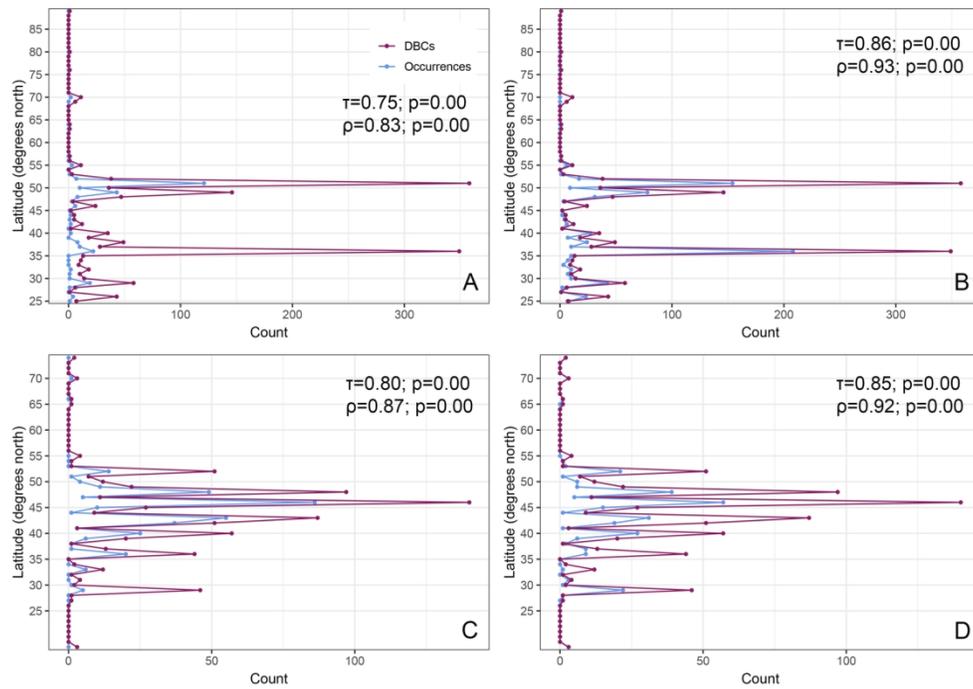


Figure 2: A, C, ceratopsid occurrences and dinosaur-bearing collections with latitude in the A, Campanian and C, Maastrichtian. B, D, hadrosaurid occurrences and dinosaur-bearing collections with latitude in the B, Campanian and D, Maastrichtian.  $\tau$  = Kendall's tau;  $\rho$  = Spearman's rho; DBCs, dinosaur-bearing collections.

148x104mm (300 x 300 DPI)

**Figure 1. A–C.** Jackknifed distributions of pBC values for Campanian (green) and Maastrichtian (purple) ceratopsids. **D.** Rarefaction curves for Campanian (green circles) and Maastrichtian (purple triangles) ceratopsids. Error bars show the 95% confidence intervals of values obtained during rarefaction.

**Figure 2: A, C,** ceratopsid occurrences and dinosaur-bearing collections with latitude in the **A,** Campanian and **C,** Maastrichtian. **B, D,** hadrosaurid occurrences and dinosaur-bearing collections with latitude in the **B,** Campanian and **D,** Maastrichtian.  $\tau$  = Kendall's tau;  $\rho$  = Spearman's rho; **DBC**s, dinosaur-bearing collections.