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Sharks That Pass In The Night:  
Using GIS to Investigate  
Competition in the Cretaceous  
Western Interior Seaway

by Bruce S. Lieberman

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**Sharks That Pass In The Night: Using GIS to Investigate Competition in the Cretaceous Western Interior Seaway**

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# 1           **Sharks That Pass In The Night: Using GIS to Investigate**

## 2           **Competition in the Cretaceous Western Interior Seaway**

3

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### 8       **Abstract**

9           One way the effects of both ecology and environment on species can be observed in the  
10       fossil record is as changes in geographic distribution and range size. The prevalence of  
11       competitive interactions and species replacements in the fossil record has long been investigated  
12       and many evolutionary perspectives, including those of Darwin, have emphasized the importance  
13       of competitive interactions that ultimately lead one species to replace another. However,  
14       evidence for such phenomena in the fossil record is not always manifest. Here we use new  
15       quantitative analytical techniques based on geographic information systems (GIS) and PaleoGIS  
16       tectonic reconstructions to consider this issue in greater detail. The abundant, well-preserved  
17       fossil marine vertebrates of the Late Cretaceous Western Interior Seaway of North America  
18       provide the component data for this study. Statistical analysis of distributional and range size  
19       changes in taxa confirms earlier ideas that the relative frequency of competitive replacement in  
20       the fossil record is limited to non-existent. It appears that typically environmental gradients  
21       played the primary role in determining species distributions, with competitive interactions  
22       playing a more minor role.

23

24

25 Key Words: competitive replacement, GIS, Western Interior Seaway, marine vertebrates

26

27 **1. Introduction**28 *(a) Historical Perspective*

29 A central question in biogeography and evolution is what causes species' distributions to  
30 wax and wane through time. Traditionally, a dominant role has been ascribed to competitive  
31 interactions between species (Darwin 1859, MacArthur and Wilson 1972, Van Valen, 1987,  
32 Vermeij 1987, Jackson and McKinney 1990, Rosenzweig and McCord 1991, Sepkoski *et al.*  
33 2000). Classic examples include the decline and replacement of brachiopods by bivalves,  
34 mammal-like reptiles by archosaurs, cyclostome bryozoans by cheilostome bryozoans,  
35 gymnosperms by angiosperms, multituberculates by rodents, and South American mammals by  
36 North American fauna; however, these cases for the most part have not been tested in detail  
37 (Benton 1987, 1996a, Rayner and Masters 1995). The theoretical importance of competition in  
38 evolution actually pre-dates Darwinian competitively driven natural selection and can be traced  
39 back to the notion of plenitude. Plenitude ascribes a fixed number of ecological niches on Earth,  
40 with rapid evolution of life to fill all available niche space. Once filled, evolution occurs in  
41 dynamic equilibrium where individual species may arise and go extinct, but patterns of global  
42 diversity remain constant (Cifelli 1981, Walker and Valentine 1984, Benton 1987, 1996b).  
43 Darwin (1859) supported this view, particularly with his famous wedge analogy, where species  
44 are akin to wedges hammered into a surface – once the surface is filled with wedges, a new  
45 wedge may only be driven in at the expense of an older wedge being driven out (Gould and  
46 Calloway 1980, Gould 1985, Benton 1996b). From this perspective, evolution occurs by a series

47 of competitive replacements through time, species' distributions are predominantly controlled by  
48 competitive interactions with contemporaries, and interspecific competition is a primary driver of  
49 macroevolution.

50 An alternative perspective is where an existing species or clade is successful until an  
51 external perturbation results in its extinction and later replacement by a new taxon. For instance,  
52 a re-examination of the diversity patterns of brachiopods and bivalves by Gould and Calloway  
53 found these clades to be as "ships that pass in the night" (Longfellow, IN: Gould and Calloway  
54 1980); a view in accord with the notion that abiotic environmental change dictates species  
55 origination and extinction patterns (Eldredge and Cracraft 1980, Vrba 1980, 1985, Cifelli 1981,  
56 Gould 1985, Masters and Rayner 1993, Benton 1996a, 2009, Barnosky 2001, Flagstad *et al.*  
57 2001, Lieberman *et al.* 2007).

58 Of course these (and other) authors acknowledge that both factors likely play some role  
59 in evolution. Thus, here we test for evidence of interspecific competition on species'  
60 distributions over macroevolutionary timescales by concentrating on identification of  
61 competitive replacements in fossil taxa using GIS. GIS-based techniques are increasingly  
62 recognized as powerful tools for investigating evolutionary patterns and processes (Rode and  
63 Lieberman 2004, Stigall and Lieberman 2006, Costa *et al.* 2008, Kozak *et al.* 2008, Butler *et al.*  
64 2010). These methods allow for quantitative measurement of distribution and range size change  
65 during specific temporal intervals. Further, GIS analyses lend themselves to statistical analysis of  
66 negative range area correlations in species pairs through time, which can be used as a proxy for  
67 evidence of competitive replacement. The focus of this analysis is a set of marine vertebrate  
68 species from the exceptionally diverse and complete record of the Late Cretaceous Western  
69 Interior Seaway of North America. This region has been the subject of palaeobiological and

70 geological study for more than a century and has been intensely sampled. Further,  
71 palaeobiological samples can be placed in a detailed stratigraphic context.

72

73 *(b) Geological Setting*

74 The Late Cretaceous covers a 35 million year period between 100-65Ma. The Earth at  
75 this time was in a greenhouse climate state with little or no polar ice (Barron 1983, Huber *et al.*  
76 2002, Spicer 2002, Everhart 2005). As a consequence of this, and higher rates of sea floor  
77 spreading, sea-level was much higher than today. In particular, central North America was  
78 covered by a shallow epicontinental sea, the Western Interior Seaway (WIS) (i.e.  $\leq 600\text{m}$  water  
79 depth) (Hattin 1982, Kauffman and Caldwell 1993, Poulsen *et al.* 2001, Everhart 2005). The  
80 WIS represents a foreland basin formed by tectonic loading and lithospheric flexure during uplift  
81 of the Rocky Mountains to the west. This basin was inundated episodically from both boreal  
82 waters extending south from the Arctic Ocean and tropical waters extending north from the  
83 proto-Atlantic/Tethys seas (Hattin 1982, Kauffman 1984, Kauffman and Caldwell 1993,  
84 Shimada *et al.* 2006). At the end of the Early Cretaceous (late Albian,  $\sim 100\text{Ma}$ ) a global sea-  
85 level low stand separated the northern and southern arms of the WIS for the last time until the  
86 late Maastrichtian ( $\sim 65\text{Ma}$ ). Cyclic sea-level changes are recorded in the WIS as three major  
87 transgressive/regressive events: the Greenhorn Cycle (late Cenomanian-Turonian), which  
88 included the sea-level high stand for the Late Cretaceous with eustatic sea-levels upwards of  
89 250m higher than today; the Niobrara Cycle (late Coniacian – early Campanian); and the  
90 Claggett/Bearpaw Cycle (Campanian – Maastrichtian) (Hattin 1982, Kauffman 1984, Kauffman  
91 and Caldwell 1993).

92 Our understanding of the Late Cretaceous WIS is based on over one hundred years of  
93 field and laboratory work by geologists, palaeoclimatologists, and palaeobiologists. As a  
94 consequence, the tectonic, environmental, and geologic history of this area is well understood  
95 and extensively palaeobiologically sampled making it an ideal region for this type of  
96 palaeobiogeographical investigation (e.g., Hancock and Kauffman 1979, Hattin 1982, Kauffman  
97 1984, Nicholls & Russell 1990, Glancy *et al.* 1993, Russell 1993, Schroder-Adams *et al.* 1996,  
98 Sageman *et al.* 1997, Schwimmer *et al.* 1997, Keller *et al.* 2004, Everhart 2001, 2005, Becker *et al.*  
99 2006, Cobban *et al.* 2006, Shimada *et al.* 2006, Ufnar *et al.* 2008). However, extensive sampling  
100 does not always equate to representative sampling; consequently, we provide various tests to  
101 assess the quality of the WIS record and its use in palaeobiogeographical analyses.

102

## 103 **2. Materials and Methods**

### 104 *(a) Data Collection*

105 A temporal and geographic occurrence database was generated for ten Late Cretaceous  
106 WIS vertebrate taxa. Taxa included three genera of shark: three species of *Ptychodus* (*P.*  
107 *anonymus*, *P. mortoni*, and *P. whipplei*), one species of *Cretoxyrhina* (*C. mantelli*), and two  
108 species of *Squalicorax* (*S. falcatus* and *S. kaupi*); as well as two genera of mosasaur (*Platecarpus*  
109 sp., and *Tylosaurus* sp.) and one teleost genus (*Xiphactinus* sp.). The taxa included in this  
110 analysis were chosen because they are common and abundant in the WIS fossil record, persist  
111 through at least three geologic stages of the Late Cretaceous, and have been well characterized  
112 taxonomically and palaeobiologically. Further, the WIS at this time had no prominent physical  
113 barriers that might have prevented interactions between taxa.

114 Data on species' geographic and stratigraphic ranges were collected through examination  
115 of museum collections, fieldwork, and survey of the literature. The following museum

116 collections were used: Natural History Museum and Biodiversity Research Center (NHM-BI,  
117 University of Kansas); Peabody Museum of Natural History (YPM, Yale University); Texas  
118 Memorial Museum (TMM, University of Texas – Austin); Sternberg Museum of Natural History  
119 (FHSM, Fort Hays State University); University of Colorado Museum (UCB, University of  
120 Colorado – Boulder); University of Nebraska State Museum (UNSM); and the Black Hills  
121 Institute (BHI, South Dakota). These museums contain important and diverse collections of WIS  
122 taxa spanning the majority of Late Cretaceous WIS geography, and taxa in these collections are  
123 well-documented geographically and stratigraphically. All museum specimens were personally  
124 examined and identification confirmed by the authors. In cases where species identifications  
125 lacked confidence, analyses were run at the generic level (e.g. *Tylosaurus*, *Platecarpus*,  
126 *Xiphactinus*). To augment information from museums, fieldwork was conducted at Late  
127 Cretaceous sites in western South Dakota and southeastern Missouri.

128       Resolution of geographic locality data was at the county-level and better, the standard  
129 level of resolution used in other GIS-based palaeobiogeographic analyses (e.g., Rode and  
130 Lieberman 2004, Hendricks *et al.* 2008, Maguire and Stigall 2009). However, most data  
131 represent even higher resolution at the 1 mi<sup>2</sup> township, range, and section. Temporal resolution  
132 was at the level of geologic stage within the Late Cretaceous and characterized by formation and  
133 member of specimen occurrence. The resulting database consists of 762 total occurrence points;  
134 the number of occurrence points per taxonomic group varies from 31 to 197 (figure 1).

135

### 136 (b) Range Reconstructions

137       Geographic locality data for each species' occurrence was georeferenced and imported  
138 into ArcGIS v.9.2 for visual representation and spatial analysis (ESRI 2006). PaleoGIS v.3.0



139 (Scotese 1998, Ross and Scotese 2000, Rothwell Group 2007) was then used to reconstruct the  
140 palaeogeography of each stage during the Late Cretaceous following the methods of Rode and  
141 Lieberman (2004) and Stigall and Lieberman (2006) (figure 2). This step ensures that  
142 distribution and range area reconstructions minimize estimation error due to tectonic contraction  
143 and expansion in the North American plate over the course of the Late Cretaceous.

144       Once PaleoGIS was used to reconstruct the geography of a particular stage, a ten  
145 kilometer buffer was applied to each specimen occurrence point. Buffering species' locality  
146 points helps control for any error in the translation from current geographic location to deep time  
147 georeferenced latitude and longitude. Additionally, buffering gives area to point occurrence data,  
148 enabling retention of these data in the analysis. ArcGIS was then used to construct least-fit  
149 polygons for each taxon at each temporal interval. The spatial analysis software available within  
150 this program was used to calculate area of each reconstructed range. Geographic range data for  
151 all taxa are provided in electronic supplementary material, table S1.

### 152 153 *(c) Identifying Competition*

154       One way competition can be observed in the fossil record is as changes in species'  
155 distribution and range size through time. Benton (1996a, b) defined "Candidate Competitive  
156 Replacements" (CCRs) as species pairs showing negatively correlated abundance and diversity  
157 patterns over time. CCRs must involve taxa with overlapping geographic and stratigraphic  
158 ranges and should also involve comparisons between taxa with similar habitat, body size, and  
159 diet. Further, all CCRs must show a distinctly "successful" taxon (the survivor) as well as a  
160 distinctly "unsuccessful" taxon, identified by range contraction and extinction within two  
161 temporal intervals after the minimum date of origin of the "successful" taxon (Benton 1996a, b).

162 This pattern can also be identified in the fossil record as negatively correlated geographic range  
163 area through time, which can be tested for statistical significance using nonparametric rank  
164 correlation in PAST v.2.01 (Hammer *et al.* 2001) (Spearman's  $\rho$  and Kendall's  $\tau$ ,  $p \leq 0.05$ );  
165 these statistical analyses were corrected for multiple comparisons using the Bonferroni  
166 correction.

167 All taxa under investigation display geographic and stratigraphic overlap. To identify  
168 CCRs, taxa with similar inferred ecotypes were compared, as taxa within the same ecotype are  
169 most likely to have interacted competitively. The taxa in this study can be divided into two  
170 general palaeoecologies: species of *Cretoxyrhina*, *Squalicorax*, *Tylosaurus*, *Platecarpus*, and  
171 *Xiphactinus* are inferred to have been pelagic predators (e.g., Russell 1967, Williamson *et al.*  
172 1993, Everhart 2005, Rothschild *et al.* 2005, Shimada and Cicimurri 2005, Becker 2006,  
173 Shimada *et al.* 2006; see Schwimmer *et al.* 1997 for additional discussions of *Squalicorax*);  
174 species of *Ptychodus* and *Rhinobatos* are inferred to have had a nekto-benthic, durophagous  
175 lifestyle (e.g., Stewart 1988, Williamson *et al.* 1993, Everhart 2005, 2007, Shimada *et al.* 2006;  
176 see Hamm 2008, 2010 for additional discussions of *Ptychodus*). Comparisons were also  
177 conducted by genus, as species within the same genus may be more likely to have the greatest  
178 degree of competitive overlap. Finally, an agnostic approach was used, and pairwise  
179 comparisons between all taxa were considered.

180

#### 181 (d) Analysis of Bias

182 There are many phenomena that can explain why one species range might increase  
183 through time while another decreases through time. In addition to competition and other  
184 processes discussed below, an incomplete fossil record could artificially produce a pattern

185 mirroring a CCR. Incompleteness of the fossil record is a potential source of bias in any  
186 palaeontological study. As previously mentioned, the Late Cretaceous WIS has been  
187 exhaustively studied for over a century and is well characterized both in terms of its geology and  
188 palaeontology. Further, it has not undergone significant tectonic modification since the Late  
189 Cretaceous. These may all partly serve to obviate the potential problems of an incomplete fossil  
190 record. Moreover, some areas within the WIS show exceptional preservation in the form of  
191 Konservat Lagerstätte; one of these, the Smoky Hill Chalk member of the Niobrara Formation  
192 spans three temporal intervals (Coniacian, Santonian, and Campanian stages) of this study  
193 (Schwimmer *et al.* 1997, Meyer and Milsom 2001, Bottjer 2002).

194         However, this does not mean that there might not be certain taphonomic factors  
195 conspiring to cloud our understanding of biogeographic patterns in these taxa over time. Because  
196 of this, three tests were used to determine if incompleteness or bias in the WIS fossil record is  
197 artifactually influencing palaeobiogeographic patterns, including those pertaining to CCRs. First,  
198 the robustness of range area reconstructions to potential outliers was tested by resampling  
199 occurrence points for each taxon. An ‘n-1’ jackknifing procedure was utilized to estimate the  
200 resampled mean range size and associated confidence bands for each taxon during each time  
201 interval (resampled data available in electronic supplementary material, table S1). This mean  
202 range area was then subjected to nonparametric rank correlation tests and the results were  
203 compared to those obtained using original range area calculations (tests on resampled data  
204 available in electronic supplementary material, tables S3 and S4, and discussed more fully  
205 below).

206         The second test compared geographic range size in each taxon to area of available Late  
207 Cretaceous sedimentary outcrop. A high percentage of overlap between the distribution of taxa

208 and available outcrop would suggest that presence/absence of Late Cretaceous geologic record  
209 may be influencing our results. The third test aimed to identify a correlation between number of  
210 data points and geographic range size for each temporal interval. In this case, if sampling bias  
211 had an effect on our range size reconstructions, a strong positive correlation between number of  
212 data points and range size would be expected.

213

### 214 **3. Results**

#### 215 *(a) Competition in the WIS*

216 Tables 1 and 2 show the results of intrageneric range area correlations and correlations by  
217 palaeoecotypes respectively; pairwise comparisons between all taxa are included in electronic  
218 supplementary material, table S2. All species did show changes in distribution and range size  
219 through time. The majority of the species comparisons showed no evidence of interspecific  
220 competition (e.g., figure 3). A complete set of geographic comparisons for all taxa considered is  
221 provided in the electronic supplementary material (figures S1–S43). Some taxa did generally  
222 show the basic biogeographic pattern predicted for a CCR (figure 4), however, when analyzed  
223 the pattern was not found to be statistically significant. Indeed, no statistically significant  
224 negative range area correlations were identified from intrageneric comparisons, within ecotype  
225 comparisons, or when all taxa were compared, after the Bonferroni correction was applied. For  
226 instance, consider that among the four possible intrageneric comparisons, only *Squalicorax*  
227 *falcatus* and *S. kaupi* is near significance using Kendall's  $\tau$  ( $\tau = -0.69007$ ,  $p = 0.0518$ ), but the  
228 correlation is not significant after a Bonferroni correction for multiple comparisons was applied  
229 (new critical p-value of  $p \leq 0.013$ ) (table 1). Thus, it appears that for these vertebrate taxa

230 evidence for candidate competitive replacements in the Cretaceous WIS is negligible to non-  
231 existent.

232

233 *(b) Analysis of Bias*

234       Geographic range estimations using this palaeobiogeographic method may be susceptible  
235 to artificial inflation by widely flung single occurrence points. In order to assess the influence of  
236 these potential outliers on our range reconstructions, and thus pertaining to the identification of  
237 statistically significant CCRs, we re-ran all the pairwise comparisons using the estimated mean  
238 geographic range calculated by jackknifing (electronic supplementary material, table S3). The  
239 results are identical: before or after correcting for multiple comparisons, no statistically  
240 significant intrageneric or within ecotype CCRs were identified; when all taxa were compared,  
241 two CCRs only appeared statistically significant before the Bonferroni correction was applied:  
242 they were no longer significant after correction for multiple comparisons. Thus, the results from  
243 analysis of the original data and the resampled data are equivalent and the data appear robust to  
244 resampling. Consequently, outliers are not likely to be playing a significant role in influencing  
245 the results.

246       To test for the effect of available outcrop area on species distributions during the Late  
247 Cretaceous, we compared species' geographic range size with area of Late Cretaceous  
248 sedimentary record; the approximate margins of the WIS for the early, middle, and late Late  
249 Cretaceous, along with the occurrence records parsed by stage, are shown in electronic  
250 supplementary material, figure S44. Taxa were shown to occupy only 4–37% of potential habitat.  
251 Because taxa are not present in all or even the majority of available outcrop area during this time

252 period, it is unlikely that the simple availability of Late Cretaceous sedimentary record is  
253 controlling the patterns of distribution and range size change observed in this analysis.

254 A correlation of number of unique geographic localities sampled with size of geographic  
255 range reconstruction for each temporal interval in this analysis is shown in table 3 (for  
256 correlation statistics using resampled means, see electronic supplementary material, table S4).  
257 The number of unique localities was used to test for sampling bias (instead of all sampled  
258 occurrences) because this reduces artificial inflation of points sampled and maximizes the  
259 potential for finding a significant correlation (thus the test is most sensitive to identifying  
260 sampling bias). None of the stages during the Late Cretaceous show significant correlations  
261 between number of data points and size of geographic range ( $p \gg 0.007$  using Bonferroni  
262 correction for multiple comparisons) except the Coniacian stage ( $p = 0.001$ ) (table 3); the same is  
263 true of the resampled data (electronic supplementary material, table S4).

264 Many (though not all) taxa have small geographic range size during the Coniacian; it is  
265 possible that this represents a bias in collection or preservation. On the other hand, this stage is  
266 the point of origin or extinction for a number of the taxa studied (e.g., *Tylosaurus* sp.,  
267 *Platecarpus* sp., *Squalicorax kaupi* originate; *Ptychodus anonymus* and *Ptychodus whipplei* go  
268 extinct). Species commonly have small geographic range size at the point of origination and  
269 extinction (particularly if speciation occurs allopatrically in small isolated populations and  
270 extinction first involves reduction to a single population). To assess the influence of this  
271 phenomenon, these taxa were removed and the correlation statistics re-run (table 3, and  
272 electronic supplementary material, table S4). Excluding taxa originating or going extinct, the  
273 number of sampled localities during the Coniacian is no longer significantly correlated with  
274 reconstructed range size in both the original and the resampled data (see table 3 and electronic

275 supplementary material, table S4). Thus the uniquely small range size of these taxa was likely  
276 causing the suggested sampling bias during this interval.

277

#### 278 **4. Discussion**

279 This study uses new techniques in quantitative biogeographic analysis to test for the role  
280 of competitive replacement in the fossil record. We focused on species' distributions in the  
281 abundant representatives of the vertebrate fauna from the Late Cretaceous WIS, specifically  
282 looking for two-taxon comparisons suggesting competitive replacement. No two-taxon  
283 comparisons showed any statistical evidence of significant, negative geographic range  
284 correlations. These results reiterate previous analyses indicating little evidence for competitive  
285 replacement (Benton 1996a, b). Further, this suggests that something other than interspecific  
286 competition plays the predominant role in influencing species distributions over  
287 macroevolutionary time scales. Such processes were most likely abiotic environmental changes,  
288 both climatic and tectonic, as these have been shown to have had a significant impact on species  
289 distributions and macroevolution at other times in the history of life (Lieberman and Eldredge  
290 1996, Lieberman 2000, Barnosky 2001, Flagstad *et al.* 2001, Rode and Lieberman 2004, Stigall  
291 and Lieberman 2006, Hendricks *et al.* 2008, Benton 2009, Gates *et al.* 2010). There could,  
292 however, also be a substantial contribution from ecological factors such as food source tracking,  
293 intraspecific interactions, etc.

294 It is worth noting that competitive replacement may be more prevalent among species  
295 that are rare and/or geographically restricted. Such cases are difficult to identify in the fossil  
296 record, and thus by necessity our study focused on more abundant and potentially more  
297 "successful" taxa from the outset. As a consequence, even though we attempted to maximize

298 recovery of CCRs by using broad definitions of palaeoecological similarity, our estimate of the  
299 frequency of CCRs is most surely an underestimate. Nonetheless, it is based on quantitative and  
300 detailed investigation of these groups, and thus the best estimate possible at present.

301         Moreover, while we believe that our analysis includes real species using a phylogenetic  
302 species concept, it is impossible to exclude the possibility that some of these species actually  
303 represent ecomorphs within a single lineage. If this were the case, then instead of identifying  
304 cases of competitive replacement between species, our analysis would be testing for intraspecific  
305 interactions occurring between co-occurring ecomorphs. The apparent non-prevalence of  
306 competitive replacement within potentially adaptive lineages then might suggest that ecomorph  
307 evolution also may not be strongly influenced by these types of competitive interactions.

308         Ultimately, this study provides little evidence that CCRs play a defining role in shaping  
309 species' distributions at the macroevolutionary scale. The driving force is instead likely to be  
310 abiotic environmental factors, such as climate and sea-level changes, that determine species  
311 distribution and range size. Other ecological factors may have been important as well, but  
312 interspecific competition does not appear to have had a major affect on macroevolutionary  
313 patterns of species in the fossil record (Vrba 1980, Nicholls & Russell 1990, Barnosky 2001,  
314 Flagstad *et al.* 2001, Benton 2009, Gates *et al.* 2010, Venditti *et al.* 2010).

315

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326

327

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542

### 543 **Figure and Table Captions**

544 **Figure 1.** Data points showing occurrence records of Late Cretaceous marine vertebrate specimens analyzed in this  
545 study. *Xiphactinus* sp. (pink), *Platecarpus* sp. (dark green), *Tylosaurus* sp. (dark blue), *Squalicorax kaupi* (orange),  
546 *Squalicorax falcatus* (red), *Rhinobatos incertus* (light green), *Ptychodus whipplei* (white), *Ptychodus mortoni* (dark  
547 gray), *Ptychodus anonymus* (light gray), *Cretoxyrhina mantelli* (yellow). Present day outcrop of Late Cretaceous  
548 sediments is also shown (brown).

549

550 **Figure 2.** Example of PaleoGIS (Rothwell Group 2007) plate tectonic reconstruction. Distribution of *Cretoxyrhina*  
551 *mantelli* (yellow), *Tylosaurus* sp. (blue). Present day outcrop of Late Cretaceous sediments is also shown (brown).  
552 (a) PaleoGIS present day tectonic configuration. (b) PaleoGIS Coniacian reconstruction (~87Ma).

553

554 **Figure 3.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
555 uncovered for the majority of two-taxon comparisons in this study. *Tylosaurus* sp. (blue) and *Platecarpus* sp. (dark  
556 green) distributions are shown during the Late Cretaceous. Late Cretaceous stages: (a) Coniacian, (b) Santonian, (c)  
557 Campanian, (d) Maastrichtian. These taxa do not show a statistically significant negative range area correlation  
558 through time and thus are not identified as CCRs. Present day outcrop of Late Cretaceous sediments is also shown  
559 (brown).

560

561 **Figure 4.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the general predicted geographic  
562 pattern of a CCR, although the negative relationship in range size is not statistically significant. *Squalicorax falcatus*  
563 (red), and *S. kaupi* (orange) distributions are shown during the Late Cretaceous. Late Cretaceous stages: (a)  
564 Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. *S. falcatus* shows stable,  
565 though dynamic, range size until the origination of *S. kaupi* in the Coniacian (c). After this time, *S. falcatus*  
566 experiences sequential decrease in range size resulting in extinction at the end Campanian (e). This example

567 illustrates a negative relationship between the range area of two ecologically similar species within the same genus,  
568 and thus could represent a competitive replacement of *S. falcatus* by *S. kaupi*. Present day outcrop of Late  
569 Cretaceous sediments is also shown (brown).

570

571 **Table 1.** Intrageneric range area correlations. A Bonferroni correction (Sokal and Rohlf 1995) for multiple  
572 comparisons indicates a critical p-value of  $p \leq 0.013$  for statistical significance.

573

574 **Table 2.** Range area correlations among species with similar palaeoecology. (a) Inferred large, pelagic (circular  
575 vertebral centra suggesting fusiform-body) predators; (b) inferred large, nekto-benthic durophagous lifestyle. A  
576 Bonferroni correction (Sokal and Rohlf 1995) for multiple comparisons indicates a critical p-value of  $p \leq 0.002$  for  
577 statistical significance.

578

579 **Table 3.** Correlation results between number of unique geographic localities sampled and reconstructed geographic  
580 range size for each stage during the Late Cretaceous. A Bonferroni correction (Sokal and Rohlf 1995) for multiple  
581 comparisons indicates a critical p-value of  $p \leq 0.007$  for statistical significance. Coniacian\* represents correlation  
582 between number of unique geographic localities and reconstructed range size after removing taxa that either  
583 originate or go extinct during this stage.

584

### 585 **Electronic Supplementary Figure and Table Captions**

586 **Figure S1.** Paleogeographic (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
587 uncovered for *Cretoxyrhina mantelli* (yellow) and *Squalicorax falcatus* (red) during the Late Cretaceous. Late  
588 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
589 of Late Cretaceous sediments is also shown (brown).

590

591 **Figure S2.** Paleogeographic (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
592 uncovered for *Cretoxyrhina mantelli* (yellow) and *Squalicorax kaupi* (red) during the Late Cretaceous. Late  
593 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian.  
594 Present day outcrop of Late Cretaceous sediments is also shown (brown).

595

596 **Figure S3.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
597 uncovered for *Cretoxyrhina mantelli* (yellow) and *Platecarpus* sp. (dark green) during the Late Cretaceous. Late  
598 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
599 of Late Cretaceous sediments is also shown (brown).

600

601 **Figure S4.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
602 uncovered for *Cretoxyrhina mantelli* (yellow) and *Tylosaurus* sp. (blue) during the Late Cretaceous. Late Cretaceous  
603 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
604 outcrop of Late Cretaceous sediments is also shown (brown).

605

606 **Figure S5.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
607 uncovered for *Cretoxyrhina mantelli* (yellow) and *Xiphactinus* sp. (pink) during the Late Cretaceous. Late  
608 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian.  
609 Present day outcrop of Late Cretaceous sediments is also shown (brown).

610

611 **Figure S6.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
612 uncovered for *Cretoxyrhina mantelli* (yellow) and *Ptychodus anonymus* (grey) during the Late Cretaceous. Late  
613 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian. Present day outcrop of Late  
614 Cretaceous sediments is also shown (brown).

615

616 **Figure S7.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
617 uncovered for *Cretoxyrhina mantelli* (yellow) and *Ptychodus mortoni* (black) during the Late Cretaceous. Late  
618 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
619 of Late Cretaceous sediments is also shown (brown).

620

621 **Figure S8.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
622 uncovered for *Cretoxyrhina mantelli* (yellow) and *Ptychodus whipplei* (white) during the Late Cretaceous. Late

623 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian. Present day outcrop of Late  
624 Cretaceous sediments is also shown (brown).

625

626 **Figure S9.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
627 uncovered for *Cretoxyrhina mantelli* (yellow) and *Rhinobatos incertus* (light green) during the Late Cretaceous.

628 Late Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day  
629 outcrop of Late Cretaceous sediments is also shown (brown).

630

631 **Figure S10.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
632 uncovered for *Squalicorax falcatus* (red) and *Platecarpus* sp. (dark green) during the Late Cretaceous. Late  
633 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
634 of Late Cretaceous sediments is also shown (brown).

635

636 **Figure S11.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
637 uncovered for *Squalicorax falcatus* (red) and *Tylosaurus* sp. (blue) during the Late Cretaceous. Late Cretaceous  
638 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
639 outcrop of Late Cretaceous sediments is also shown (brown).

640

641 **Figure S12.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
642 uncovered for *Squalicorax falcatus* (red) and *Xiphactinus* sp. (pink) during the Late Cretaceous. Late Cretaceous  
643 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
644 outcrop of Late Cretaceous sediments is also shown (brown).

645

646 **Figure S13.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
647 uncovered for *Squalicorax falcatus* (red) and *Ptychodus anonymus* (grey) during the Late Cretaceous. Late  
648 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
649 of Late Cretaceous sediments is also shown (brown).

650

651 **Figure S14.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
652 uncovered for *Squalicorax falcatus* (red) and *Ptychodus mortoni* (black) during the Late Cretaceous. Late  
653 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
654 of Late Cretaceous sediments is also shown (brown).

655  
656 **Figure S15.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
657 uncovered for *Squalicorax falcatus* (red) and *Ptychodus whipplei* (white) during the Late Cretaceous. Late  
658 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
659 of Late Cretaceous sediments is also shown (brown).

660  
661 **Figure S16.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
662 uncovered for *Squalicorax falcatus* (red) and *Rhinobatos incertus* (light green) during the Late Cretaceous. Late  
663 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
664 of Late Cretaceous sediments is also shown (brown).

665  
666 **Figure S17.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
667 uncovered for *Squalicorax kaupi* (orange) and *Platecarpus* sp. (dark green) during the Late Cretaceous. Late  
668 Cretaceous stages: (a) Coniacian, (b) Santonian, (c) Campanian, (d) Maastrichtian. Present day outcrop of Late  
669 Cretaceous sediments is also shown (brown).

670  
671 **Figure S18.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
672 uncovered for *Squalicorax kaupi* (orange) and *Tylosaurus* sp. (blue) during the Late Cretaceous. Late Cretaceous  
673 stages: (a) Coniacian, (b) Santonian, (c) Campanian, (d) Maastrichtian. Present day outcrop of Late Cretaceous  
674 sediments is also shown (brown).

675  
676 **Figure S19.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
677 uncovered for *Squalicorax kaupi* (orange) and *Xiphactinus* sp. (pink) during the Late Cretaceous. Late Cretaceous

678 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
679 outcrop of Late Cretaceous sediments is also shown (brown).

680

681 **Figure S20.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
682 uncovered for *Squalicorax kaupi* (orange) and *Ptychodus anonymus* (grey) during the Late Cretaceous. Late  
683 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian.  
684 Present day outcrop of Late Cretaceous sediments is also shown (brown).

685

686 **Figure S21.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
687 uncovered for *Squalicorax kaupi* (orange) and *Ptychodus mortoni* (black) during the Late Cretaceous. Late  
688 Cretaceous stages: (a) Turonian, (b) Coniacian, (c) Santonian, (d) Campanian, (e) Maastrichtian. Present day outcrop  
689 of Late Cretaceous sediments is also shown (brown).

690

691 **Figure S22.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
692 uncovered for *Squalicorax kaupi* (orange) and *Ptychodus whipplei* (white) during the Late Cretaceous. Late  
693 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian.  
694 Present day outcrop of Late Cretaceous sediments is also shown (brown).

695

696 **Figure S23.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
697 uncovered for *Squalicorax kaupi* (orange) and *Rhinobatos incertus* (light green) during the Late Cretaceous. Late  
698 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian.  
699 Present day outcrop of Late Cretaceous sediments is also shown (brown).

700

701 **Figure S24.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
702 uncovered for *Platecarpus* sp. (dark green) and *Xiphactinus* sp. (pink) during the Late Cretaceous. Late Cretaceous  
703 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
704 outcrop of Late Cretaceous sediments is also shown (brown).

705

706 **Figure S25.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
707 uncovered for *Platecarpus* sp. (dark green) and *Ptychodus anonymus* (grey) during the Late Cretaceous. Late  
708 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
709 of Late Cretaceous sediments is also shown (brown).

710

711 **Figure S26.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
712 uncovered for *Platecarpus* sp. (dark green) and *Ptychodus mortoni* (black) during the Late Cretaceous. Late  
713 Cretaceous stages: (a) Turonian, (b) Coniacian, (c) Santonian, (d) Campanian. Present day outcrop of Late  
714 Cretaceous sediments is also shown (brown).

715

716 **Figure S27.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
717 uncovered for *Platecarpus* sp. (dark green) and *Ptychodus whipplei* (white) during the Late Cretaceous. Late  
718 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
719 of Late Cretaceous sediments is also shown (brown).

720

721 **Figure S28.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
722 uncovered for *Platecarpus* sp. (dark green) and *Rhinobatos incertus* (light green) during the Late Cretaceous. Late  
723 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
724 of Late Cretaceous sediments is also shown (brown).

725

726 **Figure S29.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
727 uncovered for *Tylosaurus* sp. (blue) and *Xiphactinus* sp. (pink) during the Late Cretaceous. Late Cretaceous stages:  
728 (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day outcrop  
729 of Late Cretaceous sediments is also shown (brown).

730

731 **Figure S30.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
732 uncovered for *Tylosaurus* sp. (blue) and *Ptychodus anonymus* (grey) during the Late Cretaceous. Late Cretaceous

733 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
734 outcrop of Late Cretaceous sediments is also shown (brown).

735

736 **Figure S31.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
737 uncovered for *Tylosaurus* sp. (blue) and *Ptychodus mortoni* (black) during the Late Cretaceous. Late Cretaceous  
738 stages: (a) Turonian, (b) Coniacian, (c) Santonian, (d) Campanian, (e) Maastrichtian. Present day outcrop of Late  
739 Cretaceous sediments is also shown (brown).

740

741 **Figure S32.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
742 uncovered for *Tylosaurus* sp. (blue) and *Ptychodus whipplei* (white) during the Late Cretaceous. Late Cretaceous  
743 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
744 outcrop of Late Cretaceous sediments is also shown (brown).

745

746 **Figure S33.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
747 uncovered for *Tylosaurus* sp. (blue) and *Rhinobatos incertus* (light green) during the Late Cretaceous. Late  
748 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian.  
749 Present day outcrop of Late Cretaceous sediments is also shown (brown).

750

751 **Figure S34.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
752 uncovered for *Xiphactinus* sp. (pink) and *Ptychodus anonymus* (grey) during the Late Cretaceous. Late Cretaceous  
753 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
754 outcrop of Late Cretaceous sediments is also shown (brown).

755

756 **Figure S35.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
757 uncovered for *Xiphactinus* sp. (pink) and *Ptychodus mortoni* (black) during the Late Cretaceous. Late Cretaceous  
758 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
759 outcrop of Late Cretaceous sediments is also shown (brown).

760



761 **Figure S36.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
762 uncovered for *Xiphactinus* sp. (pink) and *Ptychodus whipplei* (white) during the Late Cretaceous. Late Cretaceous  
763 stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. Present day  
764 outcrop of Late Cretaceous sediments is also shown (brown).

765  
766 **Figure S37.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
767 uncovered for *Xiphactinus* sp. (pink) and *Rhinobatos incertus* (light green) during the Late Cretaceous. Late  
768 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian.  
769 Present day outcrop of Late Cretaceous sediments is also shown (brown).

770  
771 **Figure S38.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
772 uncovered for *Ptychodus anonymus* (grey) and *Ptychodus mortoni* (black) during the Late Cretaceous. Late  
773 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
774 of Late Cretaceous sediments is also shown (brown).

775  
776 **Figure S39.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
777 uncovered for *Ptychodus anonymus* (grey) and *Ptychodus whipplei* (white) during the Late Cretaceous. Late  
778 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian. Present day outcrop of Late Cretaceous sediments  
779 is also shown (brown).

780  
781 **Figure S40.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
782 uncovered for *Ptychodus anonymus* (grey) and *Rhinobatos incertus* (light green) during the Late Cretaceous. Late  
783 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
784 of Late Cretaceous sediments is also shown (brown).

785  
786 **Figure S41.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
787 uncovered for *Ptychodus mortoni* (black) and *Ptychodus whipplei* (white) during the Late Cretaceous. Late

788 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
789 of Late Cretaceous sediments is also shown (brown).

790

791 **Figure S42.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
792 uncovered for *Ptychodus mortoni* (black) and *Rhinobatos incertus* (light green) during the Late Cretaceous. Late  
793 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
794 of Late Cretaceous sediments is also shown (brown).

795

796 **Figure S43.** PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns  
797 uncovered for *Ptychodus whipplei* (white) and *Rhinobatos incertus* (light green) during the Late Cretaceous. Late  
798 Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian. Present day outcrop  
799 of Late Cretaceous sediments is also shown (brown).

800

801 **Figure S44.** PaleoGIS (Rothwell Group 2007) reconstructions showing the approximate boundaries of the WIS and  
802 occurrence records during the Late Cretaceous stages: (a) Cenomanian, (b) Coniacian, (c) Maastrichtian. Boundaries  
803 of seaway provided with assistance of Richard Mackenzie and Peg Yacobucci. Note that boundaries represent  
804 average sea-level at any stage, not necessarily high-stand. Present day outcrop of Late Cretaceous sediments is also  
805 shown (brown).

806

807 **Table S1.** PaleoGIS range area reconstructions for each taxon during each stage of the Late Cretaceous. S = range  
808 area (km), S\* = estimated mean range area calculated by jackknifing (km), SE = standard error, L1 and L2 = 95%  
809 confidence bands on S\*.

810

811 **Table S2.** Correlation results for range area analysis of all pairwise comparisons. A Bonferroni correction (Sokal  
812 and Rohlf 1995) for multiple comparisons indicates a critical p-value of  $p \leq 0.001$  for statistical significance.

813

814 **Table S3.** Correlation results for range area analysis of all pairwise comparisons using resampled mean range  
815 estimated by jackknifing procedure. A Bonferroni correction (Sokal and Rohlf 1995) for multiple comparisons  
816 indicates a critical p-value of  $p \leq 0.001$  for statistical significance.

817

818 **Table S4.** Correlation results between number of unique geographic localities sampled and reconstructed geographic  
819 range size for each stage during the Late Cretaceous using resampled mean range estimated by jackknifing  
820 procedure. A Bonferroni correction (Sokal and Rohlf 1995) for multiple comparisons indicates a critical p-value of  $p$   
821  $\leq 0.007$  for statistical significance. Coniacian\* represents the correlation between number of unique geographic  
822 localities and reconstructed range size after removing taxa that either originate or go extinct during this stage.

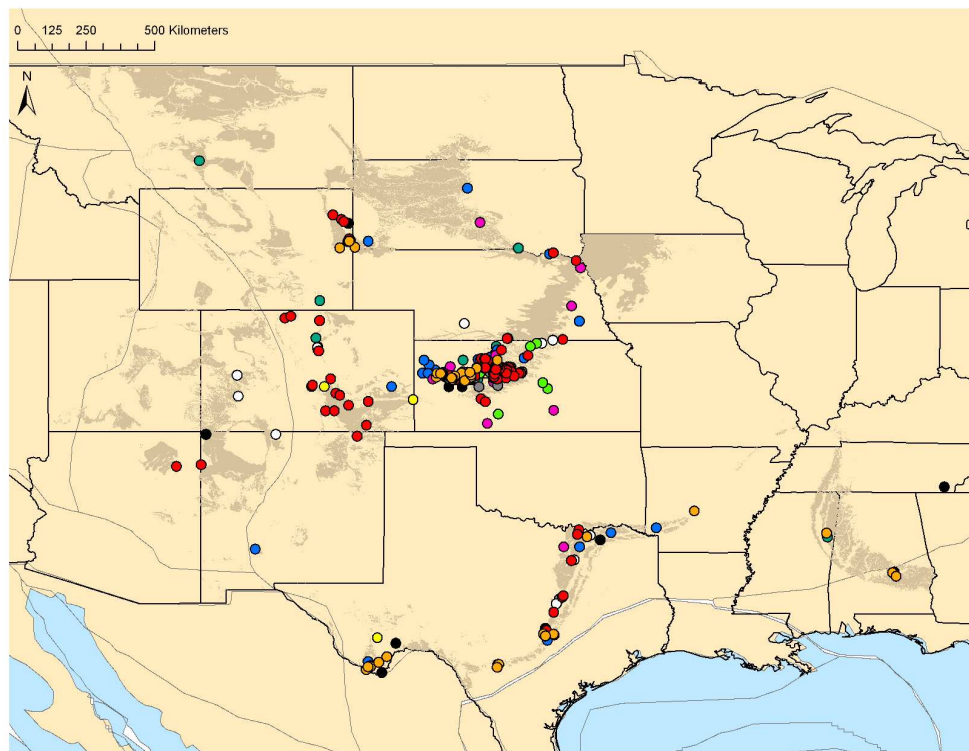


Figure 1. Data points showing occurrence records of Late Cretaceous marine vertebrate specimens analyzed in this study. *Xiphactinus* sp. (pink), *Platecarpus* sp. (dark green), *Tylosaurus* sp. (dark blue), *Squalicorax kaupi* (orange), *Squalicorax falcatus* (red), *Rhinobatos incertus* (light green), *Ptychodus whipplei* (white), *Ptychodus mortoni* (dark gray), *Ptychodus anonymus* (light gray), *Cretoxyrhina mantelli* (yellow). Present day outcrop of Late Cretaceous sediments is also shown (brown).

279x215mm (300 x 300 DPI)

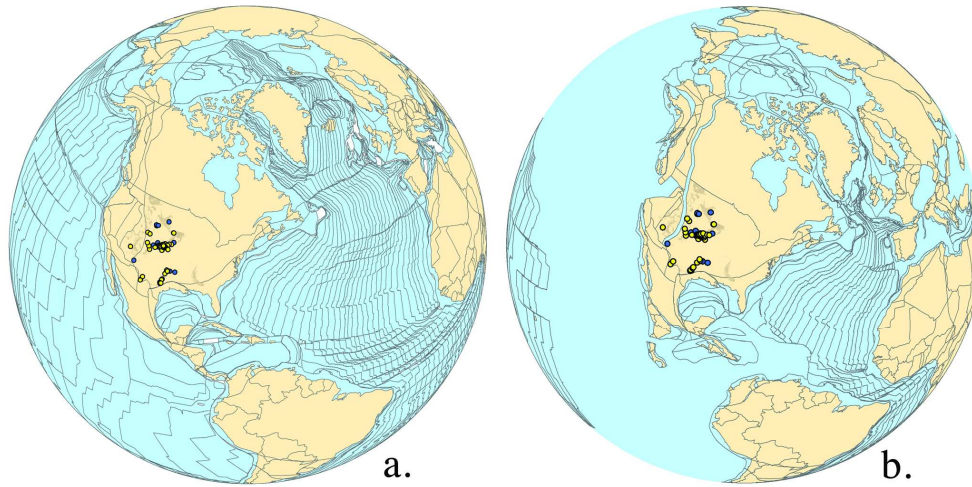


Figure 2. Example of PaleoGIS (Rothwell Group 2007) plate tectonic reconstruction. Distribution of *Cretoxyrhina mantelli* (yellow), *Tylosaurus* sp. (blue). Present day outcrop of Late Cretaceous sediments is also shown (brown). (a) PaleoGIS present day tectonic configuration. (b) PaleoGIS Coniacian reconstruction (~87Ma).  
177x87mm (300 x 300 DPI)

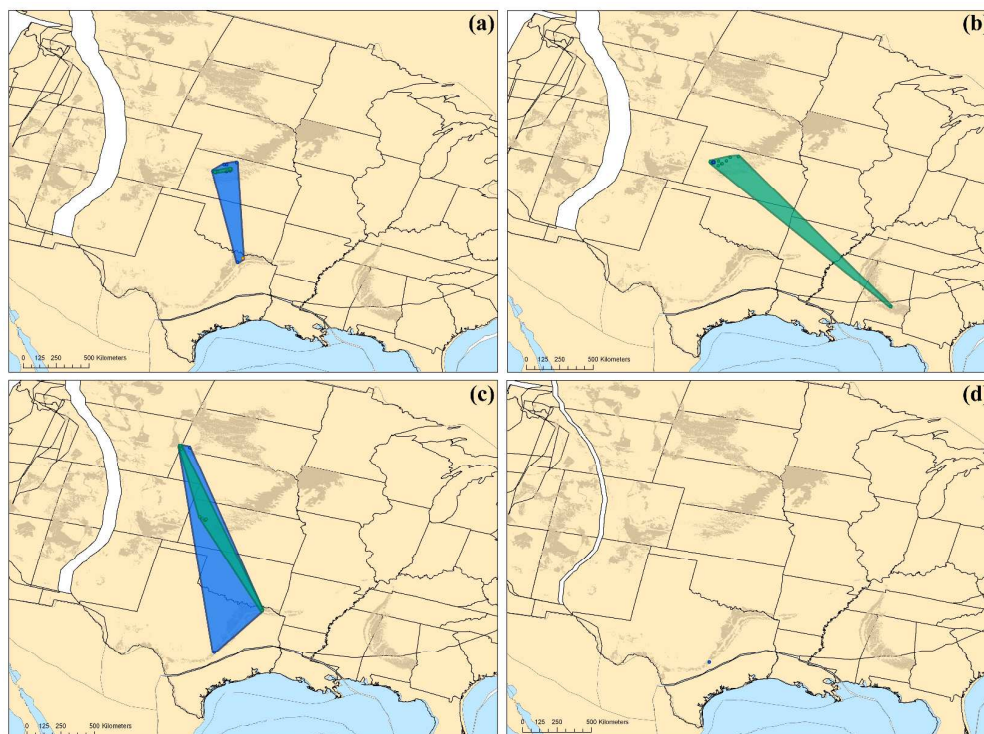


Figure 3. PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the palaeobiogeographic patterns uncovered for the majority of two-taxon comparisons in this study. *Tylosaurus* sp. (blue) and *Platecarpus* sp. (dark green) distributions are shown during the Late Cretaceous. Late Cretaceous stages: (a) Coniacian, (b) Santonian, (c) Campanian, (d) Maastrichtian. These taxa do not show a statistically significant negative range area correlation through time and thus are not identified as CCRs. Present day outcrop of Late Cretaceous sediments is also shown (brown).  
278x206mm (300 x 300 DPI)



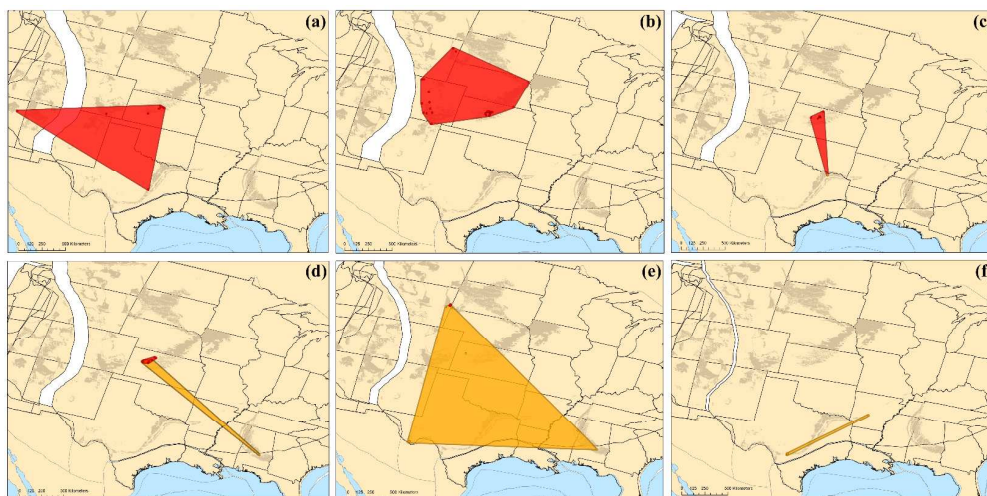


Figure 4. PaleoGIS (Rothwell Group 2007) range reconstructions illustrating the general predicted geographic pattern of a CCR, although the negative relationship in range size is not statistically significant. *Squalicorax falcatus* (red), and *S. kaupi* (orange) distributions are shown during the Late Cretaceous. Late Cretaceous stages: (a) Cenomanian, (b) Turonian, (c) Coniacian, (d) Santonian, (e) Campanian, (f) Maastrichtian. *S. falcatus* shows stable, though dynamic, range size until the origination of *S. kaupi* in the Coniacian (c). After this time, *S. falcatus* experiences sequential decrease in range size resulting in extinction at the end Campanian (e). This example illustrates a negative relationship between the range area of two ecologically similar species within the same genus, and thus could represent a competitive replacement of *S. falcatus* by *S. kaupi*. Present day outcrop of Late Cretaceous sediments is also shown (brown).

277x138mm (300 x 300 DPI)

Only

**Table 1.**

<b>Taxon A</b>	<b>Taxon B</b>	<b>Spearman's <math>\rho</math></b>	<b>p-value</b>	<b>Kendall's <math>\tau</math></b>	<b>p-value</b>
<i>Squalicorax falcatus</i>	<i>Squalicorax kaupi</i>	-0.812	0.072	-0.690	0.052
<i>Ptychodus anonymus</i>	<i>Ptychodus mortoni</i>	-0.185	0.742	-0.077	0.828
<i>Ptychodus anonymus</i>	<i>Ptychodus whipplei</i>	0.936	0.025	0.833	0.019
<i>Ptychodus mortoni</i>	<i>Ptychodus whipplei</i>	0.092	0.883	0.077	0.828

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Table 2.

Taxon A	Taxon B	Spearman's $\rho$	p-value	Kendall's $\tau$	p-value
<b>(a) Inferred pelagic, predatory taxa</b>					
<i>Cretoxyrhina mantelli</i>	<i>Squalicorax falcatus</i>	0.928	0.022	0.828	0.020
<i>Cretoxyrhina mantelli</i>	<i>Squalicorax kaupi</i>	-0.882	0.036	-0.786	0.027
<i>Cretoxyrhina mantelli</i>	<i>Tylosaurus</i> sp.	-0.765	0.097	-0.643	0.070
<i>Cretoxyrhina mantelli</i>	<i>Platecarpus</i> sp.	-0.431	0.392	-0.386	0.277
<i>Cretoxyrhina mantelli</i>	<i>Xiphactinus</i> sp.	0.174	0.733	0.138	0.697
<i>Squalicorax falcatus</i>	<i>Squalicorax kaupi</i>	-0.812	0.072	-0.690	0.052
<i>Squalicorax falcatus</i>	<i>Tylosaurus</i> sp.	-0.696	0.144	-0.552	0.120
<i>Squalicorax falcatus</i>	<i>Platecarpus</i> sp.	-0.334	0.533	-0.298	0.401
<i>Squalicorax falcatus</i>	<i>Xiphactinus</i> sp.	0.371	0.419	0.333	0.348
<i>Squalicorax kaupi</i>	<i>Tylosaurus</i> sp.	0.765	0.097	0.571	0.107
<i>Squalicorax kaupi</i>	<i>Platecarpus</i> sp.	0.770	0.108	0.617	0.082
<i>Squalicorax kaupi</i>	<i>Xiphactinus</i> sp.	0.058	0.933	0.000	1.000
<i>Platecarpus</i> sp.	<i>Tylosaurus</i> sp.	0.524	0.283	0.463	0.192
<i>Platecarpus</i> sp.	<i>Xiphactinus</i> sp.	0.152	0.833	0.149	0.674
<i>Tylosaurus</i> sp.	<i>Xiphactinus</i> sp.	-0.058	0.933	-0.138	0.697
<b>(b) inferred nekto-benthic, durophagus taxa</b>					
<i>Ptychodus anonymus</i>	<i>Ptychodus mortoni</i>	-0.185	0.7417	-0.0772	0.828
<i>Ptychodus anonymus</i>	<i>Ptychodus whipplei</i>	0.936	0.0250	0.8333	0.019
<i>Ptychodus anonymus</i>	<i>Rhinobatos incertus</i>	0.880	0.0500	0.7454	0.036
<i>Ptychodus mortoni</i>	<i>Ptychodus whipplei</i>	0.092	0.8833	0.0772	0.828
<i>Ptychodus mortoni</i>	<i>Rhinobatos incertus</i>	-0.058	0.9333	0.0000	1.000
<i>Ptychodus whipplei</i>	<i>Rhinobatos incertus</i>	0.7860	0.1167	0.5963	0.093

**Table 3.**

<b>Stage</b>	<b>Spearman's <math>\rho</math></b>	<b>p-value</b>	<b>Kendall's <math>\tau</math></b>	<b>p-value</b>
Cenomanian	0.886	0.016	0.733	0.039
Turonian	0.775	0.049	0.683	0.031
Coniacian	0.905	0.001	0.805	0.001
Coniacian*	0.700	0.2333	0.600	0.142
Santonian	0.551	0.163	0.743	0.101
Campanian	0.764	0.056	0.651	0.040
Maastrichtian	0.886	0.667	0.817	0.201
Total (combined)	0.733	0.020	0.556	0.025

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