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1 **Testing the utility of gaussian mixture models in parataxonomic**

2 **classification of fossil eggs**

3
4 *Keywords:* Gaussian mixture models; Dinosaur eggshells; Parataxonomy; Biodiversity;
5 Megaloolithidae; Dinosaur reproduction.

6 **Summary**

7 Fossil eggs and eggshells are very seldom associated with skeletal or dermal remains and can
8 rarely be assigned taxonomically. This has resulted in oologists creating an alternative
9 classification system for fossil eggs and eggshells: the parataxonomy of fossil eggs. Under
10 parataxonomy, fossil eggs are classified in oospecies, oogenera and oofamilies according to
11 qualitative characters (e.g. shape of the shell units and ornamentation) and highly heritable
12 quantitative characters (e.g. height of unit, HU, and width of unit, WU). *Megaloolithus*
13 oospecies, which have previously been partially attributed to both titanosaurian
14 and hadrosaurid dinosaurs. These oospecies are defined by are defined by the shape of their
15 units, quantification of the morphometric variation and variability of their microstructures
16 (HU and WU of eggshell units). Two competing interpretations of the observed
17 morphological variation of the eggshell units have been proposed: 1) different megaloolithic
18 morphologies are indicative of different dinosaur species; and 2) the same dinosaur species
19 was responsible for all the variation seen in megaloolithic eggshell units. In this study, a
20 Gaussian mixture model was applied to test both interpretations. This probabilistic model
21 assumes all the data points are generated from a mixture of a finite number of Gaussian
22 distributions with unknown parameters. We compared the morphometric distributions of HU
23 and WU in eggshells, eggs and clutches belonging to 7 *Megaloolithus* oospecies from
24 Catalonia to the defined morphometric variation seen in the eggshell microstructures of two
25 extant turtle species and two crocodile species. The resulting Gaussian model was best

26 defined for four distributions of HU and WU representing the extant turtle and crocodile
27 oospecies, and three distributions consisting of one or more *Megaloolithus* oospecies. Thus,
28 *Megaloolithus* oospecies from the Late Cretaceous of Catalonia depict the polymorphism of a
29 monotypic taxon and were laid by a single dinosaur species.

30 **1. Introduction**

31 Parataxonomy creates a systematic taxonomic framework for oological material that cannot
32 be directly or indirectly associated to osseous and embryonic remains *in ovo* and *in utero*,
33 eggs and eggshells *in utero*, DNA, and correlations of bones and palaeoöological material
34 from independent deposits (Varricchio *et al.* 2002; Sato *et al.* 2005; Oskam *et al.* 2010;
35 Huynen *et al.* 2010).

36 Parataxonomy defines oospecies via quantification of the polymorphism of egg
37 characters (eggshell thickness, egg size, details of ornamentation, and external pore patterns)
38 (Mikhailov 1991, 1997a, 1997b; Mikhailov *et al.*, 1996). For example, the dinosaurian
39 megaloolithid oospecies have been defined by the intra- (between eggs and clutches) and inter-
40 specific (oospecies) morphometric variability of the height and width of their units (Vianey-
41 Liaud and Lopez-Martinez, 1997; Vianey-Liaud *et al.*, 2003; Panadès I Blas, 2005).

42 Megaloolithid oospecies have been used to estimate the diversity of dinosaurs during
43 the Late Jurassic as well as the Late Cretaceous in areas where eggshell material is more
44 abundant than diagnostic skeletal remains (e.g. Zhao and Ding, 1976; Vianey-Liaud and
45 Lopez-Martinez, 1997; Vianey-Liaud *et al.*, 2003; Panadès I Blas, 2005;). However, the
46 role of megaloolithid palaeoöospecies as indicators of nesting palaeobiodiversity is debated,
47 because palaeologists disagree with the taxonomic validity of megaloolithid
48 palaeoöospecies (Vianey-Liaud and Lopez-Martinez, 1997; Peitz, 2000; Vianey-Liaud *et al.*,
49 2003; Panadès I Blas, 2005). Two hypotheses have been proposed to explain the taxonomic

50 validity of megaloolithid palaeooöpecies (Vianey-Liaud and Lopez-Martinez, 1997; Peitz,
51 2000; Vianey-Liaud *et al.*, 2003; Panadès I Blas, 2005):

52 1. Each megaloolithid palaeooöspecies is defined by dimensional variations of the
53 same eggshell's fan-shape unit, and represents dinosaur polytypic diversity. Therefore, the
54 diversity of megaloolithid palaeooöspecies reflects the diversity of dinosaur
55 palaeobiodiversity (Vianey-Liaud and Lopez-Martinez, 1997; Vianey-Liaud *et al.*, 2003;
56 Panadès I Blas, 2005).

57 2. Or, megaloolithid palaeooötaxa are characterised by different variations of the same
58 eggshell's fan-shape unit from different areas of the eggs, and describes the polymorphism of
59 a monotypic taxon, and consequently are not useful indicators of palaeobiodiversity (Peitz,
60 2000).

61 Here, we test both hypotheses using a Gaussian Mixture Model. We argued that we
62 can solve the taxonomic validity of *Megaloolithus* oospecies, by comparing the morphometric
63 variability of their eggshells to the variability of oospecies of defined layers such as modern
64 turtle and crocodile oospecies, using a Gaussian Mixture Model.

65 Gaussian mixture models predict the probability of distributions (set of points) to
66 belong to a set distribution of a classes (Reynolds, 2015). The Gaussian mixture model will
67 determine whether the morphometric variability of *Megaloolithus* oospecies is equivalent to
68 the variability of modern reptile oospecies, or no.

69

70

2. Materials

71 Megaloolithid and modern reptile eggs and eggshells, and data of megaloolithid material from
72 previous studies were used in this project. Three nests were sampled from Catalan localities: a
73 nest of two eggs from Coll de Nargó (Na) number IPS-33141, a nest of three eggs from
74 Fumanya (Fu) number IPS-27380, a nest with three eggs from Faidella (Fai) number IPS-

75 36341; and the fossil turtle nest from El Canyet number IPS-36381. The three nests are stored
76 at the Institut Català de Paleontologia Miquel Crusafont (Sabadell, Catalonia). Data of
77 megaloolithid eggshells from the Catalan localities Suterranya are from Panadès I Blas, 2005;
78 Basturs and Coll de Nargó from Peitz, 2000; and Biscarri clutch from López-Martínez *et al.*,
79 2000.

80 Reptilian nests of extant species of turtle and crocodile were used as comparative
81 material for the analysis, because the structure of their eggshells is the most similar to
82 megaloolithid palaeoocgenera (Mikhailov, 1991, 1997a, 1997b; Mikhailov *et al.*, 1996).
83 Megaloolithid, crocodile, and turtle eggshells are composed of discrete units along the
84 eggshell thickness that extend from the interior to the exterior of the thickness of the shell.

85 The eggs used were laid by the same female, and are of two different species of turtle
86 and crocodile, to capture the regional intraspecific variation of HU and WU exhibited by
87 reptile and megaloolithid oospecies (Mikhailov, 1991, 1997a, 1997b; Mikhailov *et al.*, 1996;
88 Vianey-Liaud and López Martínez, 1997). The materials used were two clutches with three
89 and two eggs of *Geochelone carbonaria* (Gc), two clutches with three eggs each of *Testudo*
90 *hermanni* (Th), donated by Bristol Zoo; four clutches with three eggs each of *Crocodylus*
91 *niloticus* (Cn) from La Ferme aux Crocodiles Zoo (France), four clutches with three eggs of
92 *C. porosus* (Cp) obtained from the Koorana Crocodile Farm (Queensland, Australia).

93 Megaloolithid and modern reptile eggs and eggshells, and data of megaloolithid
94 material from previous studies were used in this project. Three nests were sampled from
95 Catalan localities: a nest of two eggs from Coll de Nargó (Na) number IPS-33141, a nest of
96 three eggs from Fumanya (Fu) number IPS-27380, a nest with three eggs from Faidella (Fai)
97 number IPS-36341. The nests are stored at the Institut Català de Paleontologia Miquel
98 Crusafont (Sabadell, Catalonia). Data of megaloolithid eggshells from the Catalan localities

99 Suterranya (from Panadès I Blas, 2001); Basturs and Coll de Nargó (from Peitz, 2000);
100 Biscarri clutch (from López-Martínez *et al.*, 2000) were computed in the analyses.

101 Seven *Megaloolithus* oospecies were represented in the material: *Megaloolithus*
102 *baghesis* (Coll de Nargó), *M. mamillare* (Coll de Nargó, Basturs), *M. siruguei* (Biscarri, Coll
103 de Nargó, Faidella, and Fumanya), and *M. pseudomamillare*, *M. cf. mamillare*, *M. trempii*,
104 and unknown oospecies (Suterranya) (Vianey-Liaud and López Martínez, 1997; Panadès I
105 Blas, 2005; Vila *et al.*, 2010, 2011). The three nests have never been assigned
106 parataxonomically to a megaloolithid oospecies, but they exhibit the typical characters of *M.*
107 *siruguei*: subspherical eggs of 20×18.5 cm; eggshell thickness ranging between 1.75 and 3.6
108 mm; elongated fan-shaped units five times higher than wide exhibiting intermittent fusion;
109 arched growth lines running from the base to the top of the units and crossing to adjacent
110 units (Figure 1), (Vianey-Liaud *et al.*, 1994; Elez and López-Martínez, 2000; Vianey-Liaud
111 and Zelenitsky, 2003; Sellés *et al.*, 2013).

112 Reptilian nests of extant species of turtles and crocodile were used as comparative
113 material for the analysis. The eggshells units of turtles and crocodile and megaloolithid
114 eggshells exhibit a similar biomineral structure: a series of discreet units along the eggshell
115 thickness that extend from the interior to the exterior of the thickness of the shell (Mikhailov,
116 1991, 1997a, 1997b; Mikhailov *et al.*, 1996).

117 The eggs used were laid by the same female, of two different species of turtles and
118 crocodile eggs, to capture the regional intraspecific variation of HU and WU exhibited by
119 reptile and megaloolithid oospecies (Mikhailov, 1991, 1997a, 1997b; Mikhailov *et al.*, 1996;
120 López-Martínez, 1997). The materials used were two clutches with three and two eggs of *G.*
121 *carbonaria* (Gc), two clutches with three eggs each of *T. hermanni* (Th), donated by Bristol
122 Zoo; four clutches with three eggs each of *C. nilocitus* (Cn) from La Ferme aux Crocodiles

123 Zoo (France), four clutches with three eggs of *C. porosus* (Cp) obtained from the Koorana
124 Crocodile Farm (Queensland, Australia).

125 **3. Methods**

126 Parataxonomy defines oospecies via the quantification of the diversity/variability of egg
127 characters: height (HU) and width (WU) of units, egg size, details of ornamentation, and
128 external pore patterns (Mikhailov, 1991, 1997a, 1997b; Mikhailov *et al.*, 1996). Only well-
129 defined and consistent ranges of these characters should be applied for taxonomic inference
130 (Mikhailov, 1991, 1997a, 1997b; Mikhailov *et al.*, 1996). Measurements of HU and width
131 WU of units will exclusively be used here because these are the most widely used in
132 parataxonomic analysis (Mikhailov, 1991, 1997a, 1997b; Mikhailov *et al.*, 1996; Schleich *et*
133 *al.*, 1994; López-Martínez *et al.*, 2000).

134 The intraspecific variation of HU and WU in reptile and megaloolithid oospecies is
135 regional (equator and poles) in eggs and highly variable in eggs and clutches (Mikhailov,
136 1991, 1997a, 1997b; Mikhailov *et al.*, 1996; Schleich *et al.*, 1994; López-Martínez *et al.*,
137 2000). For example, Schleich *et al.* (1994) and Panadès and Patnaik (2009) found that modern
138 and fossil crocodile eggs are thicker in the equatorial region than in the poles, while López-
139 Martínez *et al.* (2000) found that a single egg in a megaloolithid nest was thicker in the poles
140 and thinner in the equator. However, none of the studies established whether morphological
141 and morphometrical changes of the eggshells units are correlated.

142 Accordingly, two *Megaloolithus* and reptile eggshells from the equator and the poles
143 from each egg were sampled, prepared in petrographic thin sections, and photographed under
144 a Polarised Light Microscope (PLM) (Carpenter, 1999; García and Vianey-Liaud, 2001a,
145 2001b; Grellet-Tinner *et al.*, 2000). Unlike the ovoidal eggs of megaloolithic layers and
146 crocodiles, the poles and equator in the spherical eggs of turtles are not distinct. Two crossing
147 circles were drawn around the middle of the eggs of turtles and the poles were marked on the

148 opposite sites of one of the circles. The two eggshells were extracted, inspected under a light
149 microscope for any pathologies. Eggshells with double layering, diagenetic alterations and/or
150 eggshells without completed units were discarded (Carpenter, 1999; García and Vianey-
151 Liaud, 2001a, 2001b; Grellet-Tinner *et al.*, 2010). Thin sections were deposited at the Museu
152 de la Conca Dellà (Catalonia).

153 HU and WU from the each eggshell units were captured digitally using Image J 1.47a
154 from photographs of calibrated petrographic thin sections (Rasband, 1997-2012). HU was
155 measured from the base to the highest point in a straight vertical line; and WU, from the
156 widest straight horizontal line of the unit under the PLM. This method is more accurate than
157 measuring the eggshells using a calliper and under S.E.M. (Scanning Electron Microscope)
158 and may produce distorted data (Mikhailov, 1991, 1997a, 1997b; Mikhailov *et al.*, 1996;
159 López-Martínez, 1997; Carpenter, 1999). The microstructures of eggshells are more
160 distinguishable under the Polarised Light Microscope (PLM), as it provides realistic
161 bidimensional visualisations of the morphometrical variations of the units under a perfect 90°
162 angle (Mikhailov, 1991, 1997a, 1997b; Mikhailov *et al.*, 1996; López-Martínez, 1997;
163 Carpenter, 1999).

164 Taking measurements of eggshells on the field using a calliper and from S.E.M
165 micrographs is imprecise. While the structure of the eggshells cannot be visualised, and
166 specimens diagenetically altered, over- and/or underlay by sediment, and pathologies could be
167 undetected; obtaining eggshells measurements is also inaccurate: the specimens are not in a
168 straight 90° angle and the protruding surfaces distort the real measurements (Mikhailov, 1991,
169 1997a, 1997b; Mikhailov *et al.*, 1996; López-Martínez, 1997; Carpenter, 1999).

170 Previous quantifications of megaloolithid oospecies were limited to small samples that
171 were taxonomically constraint, and they did not test independently the relations of the HU and
172 WU distributions (e. g. García and Vianey-Liaud, 2001a and b; Vianey-Liaud and López-

173 Martínez, 1997; Panadès I Blas, 2005). In order to investigate how the measurements might
174 be related, without the site information (eggshells, eggs, and localities), a Gaussian mixture
175 model was used to establish the most probable number of distributions of megaloolithic HU
176 and WU distributions using the taxonomically constrained samples of modern reptiles.

177 A Gaussian mixture model is a probabilistic model that assumes that data points are
178 generated from a mixture of a finite number of Gaussian distributions with unknown
179 parameters (Reynolds, 2008). Fitting the best mixture of Gaussians for a given dataset (as
180 measured by the log likelihood) results in a probability distribution of classes that can be used
181 to predict the probability (posterior) of new data points belonging to those classes (Reynolds,
182 2008). Fitting Gaussian mixture models is an example of an unsupervised learning method,
183 however, the computing required for fitting a mixture of Gaussians is exponential for the
184 number of latent Gaussian distributions, so approximate inference techniques are often used
185 (Reynolds, 2008). While this does not guarantee *the* optimal solution, models do converge
186 quickly to a local optimum (Reynolds, 2008).

187 To improve the quality it is usual to fit many of these models and choose the model
188 that best fits the data, often on the basis of log likelihood or similar approach (Reynolds,
189 2008). Here the Gaussian mixture model functions from the Netlab toolbox (Nabney and
190 Bishop, 2004) are used: these functions initialise the model using a clustering process known
191 as *k-means* and then use the expectation maximisation (EM) algorithm. EM is an iterative
192 method for obtaining maximum likelihood estimates of parameters for models that depend on
193 unobserved variables: in the present case a finite number of Gaussian distributions
194 representing the relationship between measurements irrespective of collection site (Nabney
195 and Bishop, 2004).

196 Mixture models were produced for 2 to 10 centres, each repeated 10 times. Each
197 centre represented HU and WU distributions of modern reptiles and megaloolithic eggshells.

198 From the 10 repetitions for each of the models, the best log likelihood was obtained. From
199 these, as shown in Equation 1 (Nannen, 2003, p. 14), the minimum description length (or the
200 best number of distributions accounting for the measurements), L calculated, was used to
201 choose the number of centres that best represented the data. Here, D is the data set, M_k is the
202 model, k is the number of Gaussians or centres and N is the number of points in the data set.

203

204 (1)

205

206 A between groups Kolmogorov-Smirnov test was then carried out with the
207 measurements of thickness and width as the dependent variables and centre as the factor in
208 order to investigate the difference in thickness and width for each of the centres found by the
209 mixture model. The classical one-dimensional Kolmogorov-Smirnov test is a non-parametric
210 statistic for comparing two empirical distributions, which defines the largest absolute
211 difference between the two cumulative distribution functions as a measure of disagreement.
212 Adapting this test to more than one dimension is a challenge because there are $2d-1$
213 independent ways of defining a cumulative distribution function when d dimensions are
214 involved.

215 We used a Matlab implementation (Lau, 2018) of a two dimensional Kolmogorov-
216 Smirnov test (Fasano and Franceschini, 1987; Peacock, 1983). Since we are doing multiple
217 comparisons, we need to adjust the alpha level according to the number of comparisons being
218 made. The simplest and most conservative correction is the Bonferroni correction, which is
219 satisfactory for present purposes:

220 Adjusted alpha level = alpha/n comparisons = $0.05/7 = 0.0071$

221

222

4. Results

223 Comparison of the minimum description lengths for each of the Gaussian mixture models
224 revealed that a model with seven centres provided the best description of the data, suggesting
225 that the measurements could be considered to belong to seven distinct distributions (Figure 3).
226 The regression line is a very good fit to both modern reptile and the dinosaur affirming that
227 megaloolithid and modern reptile eggshells possess similar variability distributions, but it
228 scales with the bigger scale of megaloolithid eggs.

229 In fact, the variability of egg thickness among reptile and dinosaurs is very similar:
230 both are highly variable (coefficient of variation of about 20%). The mean and variability
231 (standard deviation) of dinosaur eggs are not different to reptiles, just bigger; the regression
232 line is a fit to both the reptile and dinosaur data (Figure 2).

233 The Kolmogorov-Smirnov test revealed a significant difference for both width and
234 thickness between all the centres (Tables 2 and 3).

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238

5. Discussion

239

240 The Gaussian Mixture Model has defined 7 oospecies: 4 representing the extant turtle and
241 crocodile oospecies, and 3 representing *Megaloolithus* oospecies at the end of the Cretaceous
242 in Catalonia (Figure 3). The 3 *Megaloolithus* oospecies are considered taxonomically valid
243 because they are statistically different and between them and to the taxonomically constraint
244 samples of turtle and crocodile oospecies (see Figure 2, Tables 2 and 3). It was expected that
245 *Megaloolithus* eggs exhibit similar intra- and inter- morphometric variability to reptiles
246 because of the highly regional morphometric variation (equator and poles) of their eggshells
247 (Schleich *et al.*, 1994; López-Martínez *et al.*, 2000; Panadès and Patnaik, 2009). Accordingly,

248 we explored the two hypotheses to elucidate the taxonomic validity of *Megaloolithus*
249 parataxa.

250 The Gaussian Mixture Model defined the 7 *Megaloolithus* parataxa used in the
251 analyses as variations of the 3 *Megaloolithus* oospecies contained within them (Figure 3). The
252 Gaussian Mixture Model also grouped the same sites under the same *Megaloolithus* oospecies
253 from samples collected separately from (Figure 3). Thus, hypothesis 2 is more plausible: the 7
254 *Megaloolithus* parataxa illustrate the polymorphism of a monotypic taxon and thus the three
255 oospecies from the Late Cretaceous of Catalonia were laid by a single dinosaur species.
256 Otherwise, if *Megaloolithus* parataxa were indicating polytypic dinosaur diversity as
257 hypothesis 1 states, the Gaussian Mixture Model would have defined the 7 *Megaloolithus*
258 parataxa as seven distinct and statistically independent distributions to modern reptile
259 oospecies.

260 However, there may be a possibility that each *Megaloolithus* type was laid by two or
261 more different dinosaur species. *Crocodylus acutus* and *C. moreletii* produced identical
262 eggshells, while the species occur sympatrically and hybridise in coastal wetlands through the
263 Caribbean region (Ray *et al.*, 2004; Cedeño-Vazquez *et al.*, 2006; Cedeño-Vazquez *et al.*,
264 2008). It may be plausible that different species of titanosaur and hadrosaur produced
265 identical eggshells and interacted sympatrically along similar coastal regions. This cannot be
266 assessed as only a very few solitary osseous remains have been found from the Late
267 Cretaceous localities used here. Moreover, these have never been associated with any
268 *Megaloolithus* parataxa (Vila *et al.*, 2012; Prieto-Márquez *et al.*, 2013).

269 Furthermore, The Gaussian Mixture Model detected that the thickness of *Megaloolithus*
270 eggs significantly increase towards the end of the Cretaceous (Figure 3). Increased eggshell
271 thickness in reptiles decreases gas exchange through the eggshell membranes and thus
272 increases the chance of suffocation of the embryo (Wink *et al.*, 1994; Andrews, 2004). This

273 may have produced a high embryo mortality in *Megaloolithus* eggs, and may explain the lack
274 of embryonic remains. This can also have partially contributed to the extinction of
275 titanosaurid dinosaurs at the end of the Cretaceous in Catalonia.

276 Thickness of eggshell is determined by the resource quality or feeding behaviour of
277 animals (Schaffner and Swart, 1991; Robert and Ball, 1998; Senkoylu *et al.*, 2005). Recent
278 studies discarded that titanosaurian sauropods did not alter their dietary behaviours, food type
279 or quality of feeding resources of at the end of the Cretaceous in Catalonia (Sellés *et al.*,
280 2017). Thus, other environmental factors produced physiological alterations that affected their
281 reproductive cycle and dramatically increased the thickness of their eggshells. Further
282 research is required to ascertain the causes and possible consequences of an increase of
283 eggshell thickness in titanosaur dinosaurs at the end of the Cretaceous.

284 **6. Conclusion**

285 This study shows that Gaussian mixture models are a valid tool in parataxonomic
286 classification of fossil eggs. The Gaussian mixture model has optimally quantified the
287 polymorphism of the eggshell characters (HU and WU) of *Megaloolithus* parataxa from the
288 Late Cretaceous of Catalonia (Iberia). The model has recognised the variability of
289 taxonomically constraint samples of modern reptile oospecies and determined that previously
290 established *Megaloolithus* parataxa are variations of eggs laid by the same dinosaur species.
291 The *Megaloolithus* oospecies delimited by the model are considered taxonomically valid
292 although they cannot be assigned to a dinosaur species. Further parataxonomical analyses
293 using a Gaussian mixture model will determine the taxonomic validity of *Megaloolithus*
294 parataxa from other geographical areas.

295 **7. Software**

296 Gaussian mixture model functions from the Netlab toolbox is available from:
297 <http://www.ncrg.aston.ac.uk/netlab/>.

298

299 ImageJ is available from: <http://imagej.nih.gov/ij/>.

300

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Acknowledgments

302

Add acknowledgements if you have them

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481 Table 1. Descriptive statistics for each centre found by the Gaussian mixture model

Centre	Thickness			Width			N
	Mean	SD	SE	Mean	SD	SE	
1	0.413	0.068	0.002	0.396	0.115	0.003	1602
2	0.656	0.086	0.007	0.426	0.202	0.017	136
3	0.369	0.081	0.004	0.181	0.040	0.002	452
4	2.524	0.314	0.017	0.607	0.120	0.007	335
5	3.412	0.707	0.056	0.819	0.177	0.014	157
6	2.945	0.160	0.009	1.076	0.267	0.015	299
7	1.207	0.179	0.014	0.481	0.143	0.011	157

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Table 2. Statistical comparisons between the centres produced by the Gaussian Mixture Model under alpha/n comparisons.
KS statistic

	Centre 2	Centre 3	Centre 4	Centre 5	Centre 6	Centre 7
Centre 1	0.998	0.895	0.998	0.998	1.000	0.735
Centre 2		0.982	1.000	0.981	1.000	0.998
Centre 3			1.000	0.969	1.000	0.954
Centre 4				1.000	1.000	1.000
Centre 5					0.974	0.998
Centre 6						1.000

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512 Table 3. Statistical comparisons between the centres produced by the Gaussian Mixture

513 Model under Adjusted alpha level.

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515 P-value

	Centre 2	Centre 3	Centre 4	Centre 5	Centre 6	Centre 7
Centre 1	7.07E-121	1.46E-92	1.23E-28	0	1.15E-72	1.10E-174
Centre 2		1.25E-62	6.33E-24	1.35E-101	1.84E-48	4.84E-103
Centre 3			9.99E-24	1.22E-95	1.14E-47	1.33E-90
Centre 4				9.49E-28	4.98E-21	1.41E-27
Centre 5					5.16E-63	2.95E-236
Centre 6						1.03E-65

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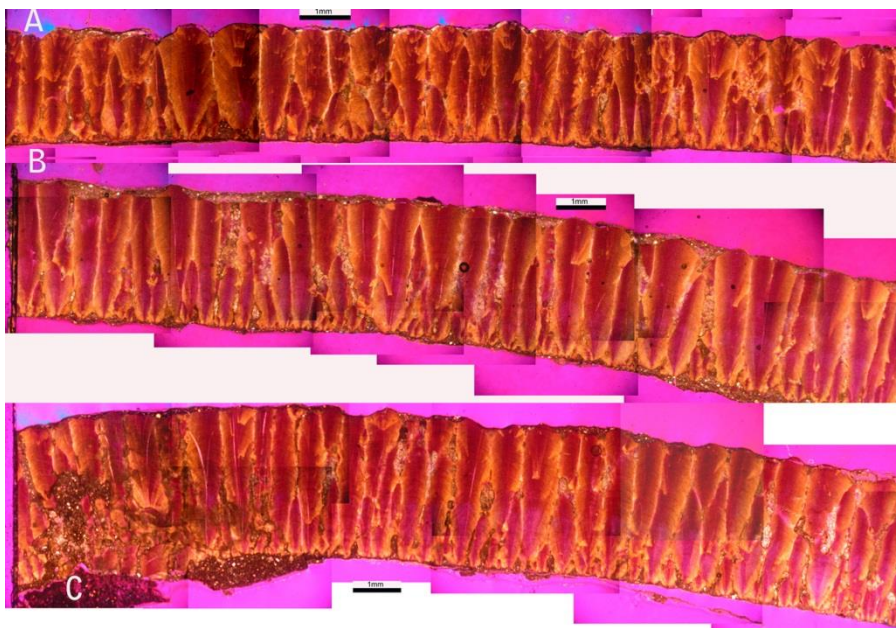
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529 Figure 1. Petrographic thin sections of eggshells from the pole 1 (A), pole 2 (B) and equator
530 (C) of an egg from the nest from Faidella used here. Notice the different morphometrical and
531 shape of units between the eggshells from the equator and the poles.

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547 Figure 2. Linear regression graphs displaying the close relation between the variation of
548 eggshell thickness between reptile and dinosaur eggs.

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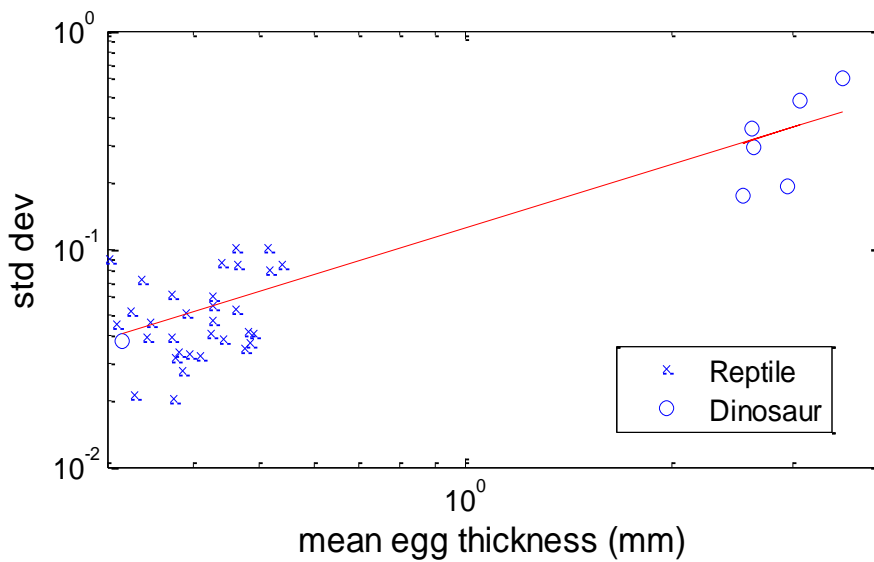
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571 Figure 3. Representation of the eggshell data together with the centres identified by the

572 Gaussian mixture model. The centres represented:

573 1 Cp = *C. porosus*

574 1 Th = *T. hermanni*

575 1 Gc = *G. carbonata*

576 1 Cn = *C. nilocitus*

577 2 St = Suterranya

578 3 Fd = Faidella

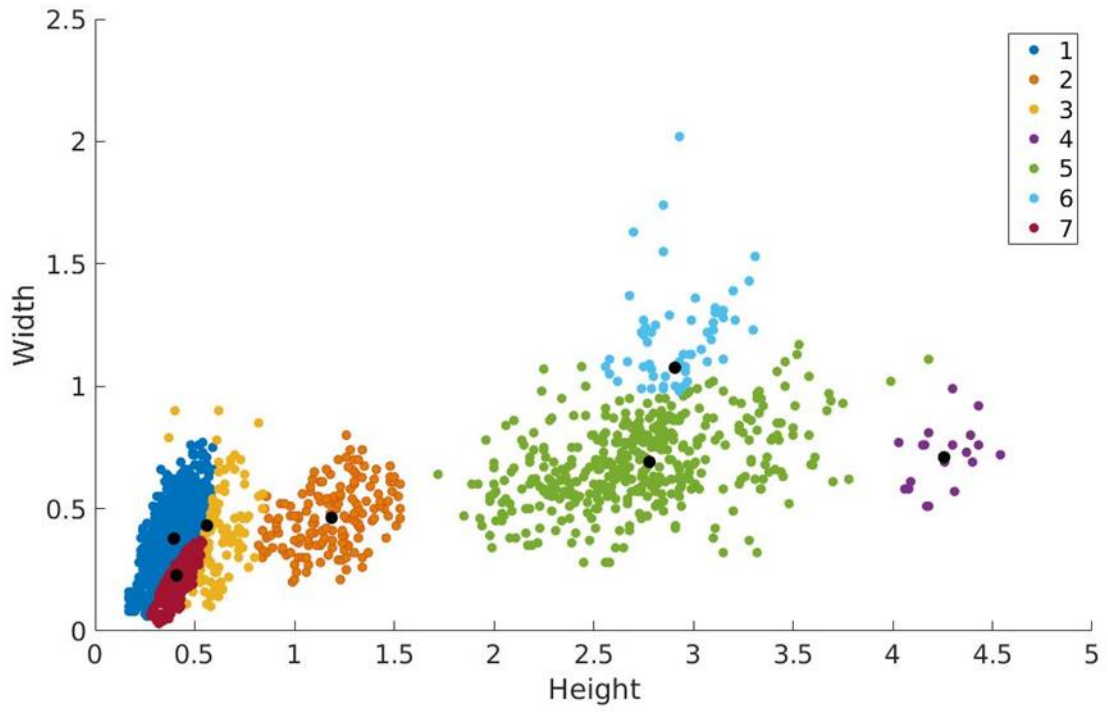
579 4 Bs = Basturs

580 5 Fd = Faidella

581 6 CdN (nest) = Coll de Nargó

582 7 CdN (locality) = Coll de Nargó.

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