1	DIVERSITY PATTERNS OF NOTOSUCHIA (CROCODYLIFORMES,
2	MESOEUCROCODYLIA) DURING THE CRETACEOUS OF GONDWANA
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13	38pp. (text+references); 5 figs.
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15	RH: POL AND LEARDI: NOTOSUCHIAN DIVERSITY PATTERNS

17 **Abstract.** Notosuchia is a diverse clade of Crocodyliformes that achieved a remarkable 18 diversity during the Cretaceous. This group is particularly abundant in continental 19 deposits of Gondwana throughout the Cretaceous, especially in South America. 20 Notosuchia was first recognized as a distinct group by the early work of Gasparini in the 21 1970's and in the last decades numerous discoveries and studies have increased the 22 geographical, temporal and taxonomical scope of this clade. Here we analyze the 23 patterns of diversity of Notosuchia during the Cretaceous, considering their taxic and 24 phylogenetic diversity, as well as implementing sampling corrections aiming to account 25 for the uneven fossil record of different stages of the Cretaceous. We identify two 26 subsequent pulses of diversification in the late Early Cretaceous and the middle Late 27 Cretaceous, followed by two separate extinction events that occurred during the latest 28 Cretaceous (Campanian/Maastrichtian). We discuss the contribution of the South 29 American, African, and Malagasy fossil records to the diversity curves, which indicates 30 the African fossil record dominates the first pulse of diversification and the South 31 American fossil record exclusively compose the second pulse of diversification. Finally, 32 we analyze the patterns of diversity shown by the different subclades of Notosuchia 33 throughout the Cretaceous, which reveal markedly different evolutionary dynamics of 34 four major groups of notosuchian crocodyliforms. 35 Key words. Notosuchia. Diversity. Radiation. Cretaceous. Gondwana. 36 **Resumen.** PATRONES DE DIVERSIDAD DE NOTOSUCHIA 37 (CROCODYLIFORMES, MESOEUCROCODYLIA) DURANTE EL CRETÁCICO 38 DE GONDWANA. Notosuchia es un clado diverso de Crocodyliformes que adquirió 39 una notable diversidad durante el Cretácico. Este grupo es particularmente abundante en 40 depósitos continentales de Gondwana a lo largo del Cretácico, especialmente en 41 América del Sur. Notosuchia fue originalmente reconocido como un grupo distintivo

42 por Gasparini en trabajos durante la década de 1970 y en años recientes numerosos 43 estudios v descubrimientos han incrementado el rango geográfico, temporal y 44 taxonómico de este clado. En este trabajo analizamos los patrones de diversidad de 45 Notosuchia durante el Cretácico, considerando su diversidad taxonómica y filogenética, 46 así como también implementando correcciones de muestreo que intentan considerar el 47 dispar registro fósil de los diferentes pisos del Cretácico. Se identifican dos pulsos 48 sucesivos de diversificación en el Cretácico Temprano tardío y en el Cretácico Tardío 49 medio, seguidos de dos eventos separados de extinción ocurridos durante el Cretácico 50 más tardío (Campaniense/Maastrichtiense). Se discuten las contribuciones del registro 51 fósil de América del Sur, África, y Madagascar a las curvas de diversidad, las cuales 52 indican que el registro africano domina el primer pulso de diversificación y el registro 53 sudamericano compone de manera exclusiva el segundo pulso de diversificación. 54 Finalmente, analizamos los patrones de diversidad de los diferentes subclados de 55 Notosuchia a través del Cretácico, los cuales revelan una dinámica evolutiva 56 marcadamente diferente para cuatro grandes grupos de crocodyliformes notosuquios. 57 Palabras clave. Notosuchia. Diversidad. Radiación. Cretácico. Gondwana. 58

60 NOTOSUCHIA is the most diverse clade of Gondwanan crocodyliforms that thrived 61 during the Cretaceous Period (Turner and Sertich, 2010; Pol et al., 2014). Much of the 62 currently known diversity, including over 70 different species, has been revealed during 63 the last 25 years through discoveries across most Gondwanan landmasses. The bulk of 64 this diversity is found in South America, where notosuchian remains have been known 65 since the latest XIXth century. The description of Notosuchus terrestris and 66 Cynodonthosuchus rothi from the Late Cretaceous of Patagonia (Woodward, 1896) was 67 the first contribution to notosuchian diversity in South America. Subsequently, during 68 the first half of the XXth century the discoveries of Uruguaysuchus (Rusconi, 1933) and 69 five taxa from the Cretaceous of Brazil (Price, 1945, 1950a,b, 1955, 1959) increased the 70 diversity of the group currently known as Notosuchia. The Brazilian taxa described by 71 Price included small-bodied forms known from relatively complete remains (i.e., 72 Araripesuchus; Price, 1959), large-bodied taxa with adaptations to hypercarnivory (i.e., 73 Baurusuchus; Price, 1945), and more fragmentary specimens with unusual tooth 74 morphology (i.e., Sphagesaurus, Itasuchus, Peirosaurus; Price, 1950a,b, 1955). 75 The systematic arrangement of these Cretaceous crocodyliforms from South 76 America was tackled in a series of influential papers by Gasparini (1971, 1972, 1981, 77 1982), who recognized and created different taxonomic groups for classifying these 78 forms. One of the most relevant contributions was the creation of Notosuchia 79 (Gasparini, 1971), a high level group that clustered small-bodied forms such as 80 Notosuchus, Araripesuchus, and Uruguaysuchus. Within Notosuchia, Gasparini (1971) 81 restricted Notosuchidae to Notosuchus and grouped the latter two species by erecting 82 the family Uruguaysuchidae. Other contributions of Gasparini were focused on less 83 diverse groups at that time, such as Sebecosuchia (Gasparini, 1972) and Peirosauridae 84 (Gasparini, 1982), which were not considered part of Notosuchia until recently (Ortega

85 et al., 2000; Pol, 2003; Turner and Sertich, 2010; Pol et al., 2012, 2014). Within 86 Sebecosuchia, Gasparini validated the Cretaceous group Baurusuchidae (including 87 Baurusuchus and the fragmentary Cynodontosuchus) as related to the Cenozoic 88 sebecids. Finally, Gasparini erected Peirosauridae (Gasparini, 1982) by recognizing the 89 distinctness of Peirosaurus torminni from all other crocodyliforms, while studying the 90 type materials of this taxon described originally by Price (1955), and more complete 91 material discovered in Patagonia. These studies therefore organized the known diversity 92 of Cretaceous crocodyliforms from South America by recognizing four basic groups: 93 Notosuchidae, Uruguaysuchidae, Peirosauridae, and Baurusuchidae. These four groups, 94 recognized by Gasparini on the basis of only six different species, represent adaptive 95 morphs that differ from each other in a suite of characters of the rostral region, 96 dentition, and palatal anatomy (Fig. 1).

97 The diversity of Cretaceous crocodyliforms from South America (and other 98 regions of Gondwana) has remarkably increased since the original studies of Gasparini 99 in the 1970's and 1980's (Carvalho and Bertini, 1999; Ortega et al., 2000; Campos et 100 al., 2001; Carvalho et al., 2004, 2005, 2007, 2011; Pol and Apesteguía, 2005; Nobre 101 and Carvalho, 2006; Andrade and Bertini, 2008; Iori and Carvalho, 2009, 2011; Kellner 102 et al., 2009, 2011a,b; Marinho and Carvalho, 2009; Novas et al., 2009; Nascimento and 103 Zaher, 2010; Montefeltro et al., 2011; Martinelli et al., 2012; Marinho et al., 2013). The 104 new discoveries, coupled with the inclusion of both Baurusuchidae and Peirosauridae 105 within Notosuchia in recent phylogenetic studies (e.g., Turner and Sertich, 2010; Pol et 106 al., 2012, 2014), revealed a previously unsuspected diversity of Notosuchian 107 crocodyliforms. There has been an almost 4-fold increase in its known diversity during 108 the last 25 years (Fig. 2). This increase in notosuchian diversity improved our 109 knowledge on the distribution of this clade, both temporally and geographically. The

biochron of most notosuchian clades is restricted to the Cretaceous, and ranges from the
Aptian (Early Cretaceous) to the Maastrichtian (Late Cretaceous), although a lineage of
notosuchians of debated affinities (Sebecidae) is recorded after the K/Pg extinction
event, and survived until the Miocene in South America (Gasparini, 1972, 1996;

114 Buffetaut, 1982; Kellner *et al.*, 2014).

115 From a biogeographic point of view, recent discoveries are highlighting a 116 growing number of Cretaceous notosuchians in other Gondwanan landmasses in 117 addition to South America, such as Africa (Sereno and Larsson, 2009; O'Connor et al., 118 2010; Sertich and O'Connor, 2014) and Madagascar (Buckley and Brochu, 1999; 119 Buckley et al., 2000; Simons and Buckley, 2009). Furthermore, fragmentary (but still 120 informative) remains found in Indo-Pakistan (Wilson et al., 2001; Prasad and de Broin, 121 2002; Prasad et al., 2013), Central Asia (Chimaerasuchus; Wu and Sues, 1996), and 122 Europe (Company et al., 2005; Dalla Vechia and Cau, 2011; Rabi and Sebök, 2015) 123 suggests that the geographic distribution of Notosuchia was broader than previously 124 thought. The major diversity of Cretaceous notosuchians is, however, still found in 125 South America, where over 70% of the known species of this group have been 126 recovered (Fig. 2).

127 These discoveries prompted a growing number of research efforts focused on the 128 anatomy and systematics of notosuchian crocodyliforms. A strong component of recent

research on notosuchians has been the use of cladistics analysis for testing the

relationships of Notosuchia (Ortega et al., 2000; Pol, 2003; Carvalho et al., 2004;

131 Turner and Sertich, 2010; Andrade et al., 2011; Montefeltro et al., 2011; Pol et al.,

132 2012, 2014). Many of these studies are now corroborating the monophyly of the four

133 Cretaceous clades early recognized by Gasparini for South American forms in pioneer

134 contributions during her early career (*i.e.*, Uruguaysuchidae, Peirosauridae,

135 Baurusuchidae, and a clade allying *Notosuchus* and closely related forms; Gasparini,

136 1971, 1972, 1982). In this contribution, we analyze the diversity patterns among

137 notosuchians during the Cretaceous, aiming to evaluate with the currently known

diversity, the radiation and extinction events of this diverse group of crocodyliforms.

### 139 MATERIALS AND METHODS

140 Notosuchian diversity

141 *Notosuchian species.* A list of published species of notosuchian crocodyliforms was

142 compiled summarizing the diversity and distribution of the group, including age,

143 geographic and stratigraphic provenance, systematic assignment to five different

subgroups of Notosuchia (see below), and year of publication (see Supplementary

145 Information).

146 Notosuchian phylogeny. The systematic arrangement of notosuchian species was based

147 on recent phylogenetic analyses published by Pol *et al.* (2014), with subsequent addition

148 of two taxa made by Leardi et al. (2015). These studies were chosen as they are the

149 most comprehensive analyses, in terms of both taxon and character sampling, of

150 notosuchian crocodyliforms performed to date and agree in many aspects to other

studies published in recent years (Sereno and Larsson, 2009; Turner and Sertich, 2010;

152 Andrade *et al.*, 2011; Montefeltro *et al.*, 2013; Sertich and O'Connor, 2014). The

topologies from these phylogenetic analyses were used for two main purposes. Firstly,

the topology served for establishing the clade assignment for each terminal taxon into

155 five major clades recognized within Notosuchia during the Cretaceous (see below).

156 Secondly, the topologies served to perform a phylogenetic correction of diversity based

157 on the inferences of ghost lineages (Norell, 1992) at each period of time (see below).

158 Some notosuchian species were not included in the data matrices published by Pol *et al.* 

159 (2014) and Leardi et al. (2015) and therefore have been excluded from the diversity

analyses conducted here, because their absence in the used phylogenetic hypotheses

161 precluded assessing their impact on the phylogenetically corrected diversity measures.

162 These, however, are limited to 18 out of the 77 known species, may not alter the

163 diversity trends discussed in this paper.

164 *Notosuchian clades.* Most recent phylogenetic analyses have recovered a large clade 165 clustering most Cretaceous crocodyliforms from Gondwana (Turner and Sertich, 2010; 166 Andrade et al., 2011; Montefeltro et al., 2013; Pol et al., 2012, 2014; Sertich and 167 O'Connor, 2014), creating a relatively recent but broad consensus on the monophyly of 168 Notosuchia and its taxonomic content. The taxonomic content of Notosuchia has been 169 enlarged in comparison with the one original proposal by Gasparini (1971; restricted to 170 Notosuchidae+Uruguaysuchidae) and currently includes the Cenozoic Sebecidae and 171 the Cretaceous Peirosauridae (and related forms; Turner and Sertich, 2010; Pol et al., 172 2012). Although some discrepancies still exist among published phylogenies (e.g., the 173 exclusion of Peirosauridae from Notosuchia), we have based our study on the 174 phylogenetic results of Pol et al. (2014) and for the purpose of assessing diversity 175 patterns we recognize here five major groups of Cretaceous notosuchians (Fig. 3), four 176 of which represent monophyletic clades. These four clades correspond to the four basic 177 taxonomic groups originally identified by Gasparini (although with some differences in 178 their taxonomic content).

The first of them, Uruguaysuchidae (Fig. 3), includes *Uruguaysuchus*, the now highly diverse *Araripesuchus* (including six species distributed from the Albian to the Campanian–Maastrichtian; Price, 1959; Buffetaut, 1981; Ortega *et al.*, 2000; Pol and Apesteguía, 2005; Turner, 2006), as well as the bizarrely broad snouted *Anatosuchus* (Sereno *et al.*, 2003). The second clade, Peirosauridae, includes a variety of Late Cretaceous South American taxa (*e.g.*, *Peirosaurus*, *Lomasuchus*, *Gasparinisuchus*,

185 Uberabasuchus, Montealtosuchus; Price, 1955; Gasparini et al., 1991; Carvalho et al., 186 2004, 2007; Martinelli et al., 2012) but also closely related African species from the 187 "mid" Cretaceous (e.g., Hamadasuchus, Stolokrosuchus; Larsson and Gado, 2000; 188 Larsson and Sues, 2007). We have included in this group the bizarre but closely related 189 clade Mahajangasuchidae (Fig. 3), recorded in the Cenomanian of Africa and the 190 Campanian–Maastrichtian of Madagascar (Sereno and Larsson, 2009). The third group, 191 here referred as basal ziphosuchians, does not represent a clade but a number of taxa 192 mainly recorded in the "middle" Cretaceous of Africa (Libycosuchus, Malawisuchus, 193 Pakasuchus; Stromer, 1914; Gomani, 1997; O'Connor et al., 2010), South America 194 (Candidodon; Carvalho, 1994), and some relictual forms in the Late Cretaceous 195 (Neuquensuchus, Simosuchus; Buckley et al., 2000; Fiorelli and Calvo, 2007). These 196 forms are clearly closer to advanced notosuchians and sebecosuchians than to 197 uruguaysuchids and peirosaurids but are placed basally within Ziphosuchia. The fourth 198 clade, advanced notosuchians (sensu Pol et al., 2014), is recorded in the Late 199 Cretaceous of South America and includes its basal members (Morrinhosuchus, 200 Notosuchus, Mariliasuchus; Woodward, 1896; Carvalho and Bertini, 1999; Iori and 201 Carvalho, 2009) and the diverse Sphagesauridae (Fig. 3). The fifth clade, 202 Baurusuchidae, is mainly recorded in two formations from the middle Late Cretaceous 203 of Argentina (Cvnodonthosuchus, Wargosuchus; Woodward, 1896; Martinelli and Pais, 204 2008) and Brazil (Baurusuchus, Campinasuchus, Pissarrachampsa, Aplestosuchus, 205 Gondwanasuchus; Price, 1945; Carvalho et al., 2011; Montefeltro et al., 2011; Godoy 206 et al., 2014; Marinho et al., 2013), with the possible addition of Pabwehshi (Wilson et 207 al., 2001) a fragmentary form found in the Maastrichtian of Pakistan. Finally, although 208 our focus is centered in the Cretaceous, we have grouped the Cenozoic Sebecidae and

allies from other regions (e.g., Iberosuchus, Bergisuchus, Doratodon; Kuhn, 1968;

Antunes, 1975; Company *et al.*, 2005; Rabi and Sebök, 2015) in a separate group.

#### 211 Diversity analysis

212 The diversity analyses performed here are based on establishing the number of 213 notosuchian lineages present at different periods of time during the Cretaceous. The 214 most basic and raw estimate is the taxic diversity (Levinton, 1988) at a given period of 215 time (*i.e.*, number of species known from that time slice). Different corrections that aim 216 to minimize the bias introduced by the vagaries of the fossil record have been proposed 217 for establishing diversity patterns across time. Here we have explored the use of two 218 commonly used corrections, one that aim to minimize the influence of the uneven 219 distribution of fossiliferous rocks for different periods of time (sampling correction) and 220 other that corrects diversity counts based on lineages inferred from the phylogenetic 221 analysis (phylogenetic correction; Norell and Novacek, 1992a,b). 222 *Time bins.* Diversity curves are not only affected by the granularity of the taxonomy 223 (hierarchical taxonomic level) but also by the precision of age assignment for each 224 fossil species. This depends on the nature of the available chronostratigraphic 225 information as well as the questions being tested. Here we have chosen to assign ages 226 based on the geological stages recognized by the International Chronostratigraphic 227 Chart (Cohen *et al.*, 2012). This follows the procedures of recent studies on the 228 diversity dynamics of mesozoic vertebrates (Barrett et al., 2009; Butler et al., 2009; 229 Mannion et al., 2011) and is based on the fact that the age of most Cretaceous 230 continental units from Gondwana is only constrained to one geological stage, at best. 231 Several lithostratigraphic units have uncertain age assignment and assigned to various 232 contiguous stages (e.g., Aptian–Albian, Turonian–Santonian). These diversity counts

233 for these units have been counted for the stages included in the uncertainty range rather

than choosing an arbitrary midpoint age assignment. These may create plateaus in the
diversity curves that should be interpreted as chronostratigraphic uncertainty rather than
stasis of diversity levels though time.

237 *Sampling correction.* The geological record heavily influences our perception of the fossil diversity, in particular limiting the findings of the taxa of interest. It has been 238 239 demonstrated that the amount of taxa known for a particular time bin is correlated with 240 the number of fossil-bearing formations available (e.g., Behrensmeyer et al., 2000; 241 Miller, 2000; Alroy et al., 2001, 2008). A typical correction used in diversity studies 242 implies the recognition of the fossil-bearing formations for the particular case of study 243 (e.g., Barrett et al., 2009; Butler et al., 2009; Irmis, 2011; Mannion et al., 2011), and 244 afterwards normalizing the diversity curves. In our case, we recognized the 245 notosuchian-bearing formations (NBFs) for each time bin (see Supplementary

Information), and the resulting diversity for that period of time was divided by thenumber NBFs.

248 *Phylogenetic correction.* The number of species known for each period of time (taxic 249 diversity) is a minimum estimate on the true diversity due to the incompleteness of the 250 fossil record. Phylogenetic trees provide hypotheses on the existence of lineages prior to 251 their first appearance in the fossil record (ghost lineages sensu Norell, 1992). A ghost 252 lineage extending throughout a period of time implies an undetected lineage that can be 253 added to the known (taxic) diversity, which is the basic rationale of the phylogenetic 254 correction of diversity curves (Norell and Novacek, 1992a,b). Here we have counted an 255 additional lineage to the diversity counts when the phylogenetic topology implied a 256 lineage the spans at least throughout an entire geological stage.

257 **RESULTS** 

258 The diversity of notosuchians varied significantly along the Cretaceous. We

show first the patterns of overall diversity through time for the clade Notosuchia.

260 Afterwards, we decompose the curves to evaluate the contribution of the different

261 geographic areas and phylogenetic clades to the total diversity of Notosuchia.

## 262 Notosuchian diversity curves

263 *Taxic and phylogenetic diversity.* The number of notosuchian species known for each 264 period of time increases in two distinct pulses during the Cretaceous. The first pulse 265 (Fig. 4.1) represents the Aptian radiation of basal notosuchians (see Pol et al., 2014). 266 The notosuchian diversity remains stable during the Aptian–Cenomanian, a stasis that is 267 likely affected by the chronostratigraphic uncertainty of notosuchian bearing formations 268 of the "middle" Cretaceous of South America (e.g., Itapecurú, Guichón) or Africa (Kem 269 Kem, Gadoufaua, Galula). After this stasis, the second pulse of diversification (Fig. 4.1) 270 occurs later during the Late Cretaceous, presumably between the Turonian and 271 Santonian, where the highest peak of notosuchian diversity is achieved (Turonian-272 Santonian radiation sensu Pol et al., 2014). The precise timing of this radiation depends 273 on the age of certain formations from South America (e.g., Adamantina, Bajo de la 274 Carpa; see Pol et al., 2014 and Discussion below). Drastic drops in diversity are 275 recorded at the end Cretaceous (most likely Campanian-Maastrichtian) that reduced the 276 diversity of Cretaceous notosuchians, of which only one lineage (sebecids) survives the 277 K/Pg extinction event.

The phylogenetic correction shows that the periods of time in which there are more unsampled lineages are precisely the timespan during which most notosuchians are known (Aptian–Santonian; light grey curve in Fig. 4.1). This indicates that both the Aptian and the Turonian–Santonian radiation events were larger than what the direct reading of the fossil record (*i.e.*, taxic diversity) implies. In our analysis, the

phylogenetic correction almost does not make any difference before the Aptian or after
the Santonian, implying that our phylogenetic tree does not indicate that unsampled
lineages are biasing the estimates of low notosuchian diversity outside the Aptian–
Santonian timespan.

287 *Sampling correction.* The sampling correction provides an overall similar pattern of 288 diversity, with two pulses of diversification (Aptian and Turonian-Santonian) and a 289 drastic drop in diversity at the latest Cretaceous (Fig. 4.2). However, there are two most 290 notable differences with respect to the diversity curves uncorrected for uneven 291 sampling. The first of them is that the first diversification pulse reaches a peak by the 292 Aptian but then is followed by a decrease in diversity (corrected by sampling) during 293 the Albian and Cenomanian (Fig. 4.2) rather than a diversity stasis. This drop is caused 294 by the fact that there are a similar number of notosuchian taxa (or lineages) known for 295 these three stages, but there are more notosuchian bearing formations for the 296 Cenomanian (nine) than for the Aptian (seven) so that the diversity relative to the 297 number of sampling units is lower. The second difference is found in the diversity 298 dynamics across the K/Pg mass extinction event. When the diversity is uncorrected by 299 sampling there is a drastic drop in diversity between the Maastrichtian and the 300 Paleocene (Fig. 4.1) whereas there is only a minor decrease in diversity across the K/Pg 301 boundary when the diversity is corrected by uneven sampling (Fig. 4.2). The end 302 Cretaceous known diversity is indeed higher than in the Paleocene (11 species versus 4 303 species) but the number of units for these two periods of time is also markedly different. 304 The apparent insensitivity of notosuchians to the mass extinction event inferred from 305 the diversity curve corrected by sampling is nonetheless a byproduct of lumping all 306 notosuchians within a single taxonomical category (see below).

307 Diversity patterns of Notosuchia across Gondwana

As noted above, although close to 75% of the known notosuchian diversity is found in South America (Fig. 2), there is a large number of notosuchian species known from Africa, Madagascar, and other regions of the world. The diversity of Notosuchia varies not only across time but also across space. The geographically split diversity curves aim to show the different contributions of Gondwanan landmasses during the Cretaceous (Fig. 5.1).

314 The first diversification event (Aptian radiation) is largely formed by the African 315 fossil record (Fig. 5.1), which accounts for more than half the known notosuchian 316 diversity in the Early Cretaceous. The large African contribution to this radiation is 317 surely influenced by the numerous notosuchians discovered in highly fossiliferous 318 deposits of Niger (Sereno and Larsson, 2009) but also in other regions of Africa (e.g., 319 Tanzania, Malawi, Egypt, Morocco; Stromer, 1914; Gomani, 1997; Sereno and Larsson, 320 2009; O'Connor et al., 2010; Sertich and O'Connor, 2014). The contribution of the 321 South American fossil record to the first diversification pulse is second to the African 322 contribution and increases towards the Cenomanian. The record in other landmasses is 323 limited to the Aptian-Albian Chimaerasuchus paradoxus from China (Wu et al., 1995). 324 The second diversification pulse (referred as the Turonian–Santonian radiation) 325 is exclusively formed by the South American fossil record (Fig. 5.1), which accounts 326 for the major diversity peak in the history of Notosuchia. This is undoubtedly 327 influenced by the remarkable diversity of notosuchians in the Adamantina Formation in 328 Brazil and the Neuquén Group in northwestern Patagonia (see Pol et al., 2014), which 329 represents a major radiation not only in terms of the number of taxa but also in the 330 ecological diversity of the group known from these units (Godoy et al., 2014). As noted 331 above, some authors regard the age of this unit as younger than Turonian–Santonian

(e.g., Gobbo-Rodrigues et al., 1999; Fernandes and Coimbra, 2000) and therefore this 332 333 diversity peak may be displaced slightly later, towards the Campanian (see below). 334 By the latest Cretaceous (Campanian–Maastrichtian), notosuchian diversity falls 335 to half the number of species known from the preceding diversity peak. Although the 336 decrease in South American diversity is steep, this landmass still contributes to more 337 than half of the known diversity in the latest Cretaceous (Fig. 5.1). An important 338 contribution to the Campanian–Maastrichtian diversity is found in Madagascar 339 (Maevarano Formation; Buckley and Brochu, 1999; Buckley et al., 2000; Turner, 2006; 340 Rasmusson Simons and Buckley, 2009). Other regions have provided records of 341 ziphodont crocodyliforms by the end Cretaceous linked to different groups of 342 notosuchians, such as Europe (Company et al., 2005) or Pakistan (Wilson et al., 2001). 343 After the K/Pg extinction event the diversity of notosuchians is mainly restricted to 344 South America, with minor contributions from other regions (Fig. 5.1), such as Europe 345 (e.g., Iberosuchus, Bergisuchus; Antunes, 1975; Ortega et al., 1996; Ortega, 2004) 346 although possible notosuchian remains have also been reported from the Paleogene of 347 Africa (Eremosuchus; Buffetaut, 1989).

# 348 Diversity patterns of notosuchian subclades

349 The diversity of Notosuchia as a whole varies markedly during the Cretaceous, 350 and this variation responds to the sum of evolutionary dynamics of the different 351 notosuchian subclades. The taxic diversity curves of the five major groups of 352 Cretaceous notosuchians reveal that different clades diversify and become extinct at 353 different times (Fig. 5.2). Three of the five groups (basal members of Ziphosuchia, 354 uruguaysuchids, and peirosaurids) show a similar pattern of taxic diversity: they early 355 diversify and form the first pulse of diversification during the "middle" Cretaceous 356 (Aptian–Cenomanian), followed by a period of low diversity in the Turonian–

357 Santonian, a minor diversity peak during the latest Cretaceous (Campanian-

358 Maastrichtian), and a complete extinction by the end of the Mesozoic (Fig. 5.2). The

two other groups (advanced notosuchians and baurusuchids) differ markedly from this

360 pattern of diversification, but their diversity curves closely resemble to each other.

361 These two groups form the major diversification peak of Notosuchia (Turonian-

362 Santonian) but have extremely low taxic diversity before and after this period.

363 Uruguaysuchids have an initial diversity represented by the *Araripesuchus* 

364 species from South America and Africa and the African *Anatosuchus*. This group

365 apparently becomes extinct throughout Gondwana, except for Madagascar, where a

366 lineage of *Araripesuchus* survived until the latest Cretaceous (*A. tsangatsangana*;

367 Turner, 2006). The age of *Uruguaysuchus* from South America, however, is uncertain

368 (Soto et al., 2011) and precludes defining the mid-Cretaceous extinction of

369 uruguaysuchids in South America more confidently.

370 The basal ziphosuchians follow a similar pattern of diversity as uruguaysuchids

371 (Fig. 5.2), having a high initial diversification that is broadly distributed, but dominated

372 by the "middle" Cretaceous African record (*Libycosuchus*, *Pakasuchus*, *Malawisuchus*).

373 Subsequently basal ziphosuchians become extinct, except in Madagascar where a

374 surviving lineage is represented by the aberrant *Simosuchus* in the latest Cretaceous

375 (Buckley *et al.*, 2000; Turner and Sertich, 2010).

376 Peirosaurids (and closely related forms) also have their initial diversity in the377 Aptian and through the "middle" Cretaceous, being recorded in South America

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378 (Barcinosuchus; Leardi and Pol, 2009) and Africa (Hamadasuchus, Stolokrosuchus,

379 Kaprosuchus; Larsson and Gado, 2000; Larsson and Sues, 2007; Sereno and Larsson,

380 2009). However, they differ slightly from the two previous groups in that they are

381 recorded in South America throughout the rest of the Cretaceous, during the Turonian-

382 Santonian (Lomasuchus, Gasparinisuchus, Montealtosuchus; Gasparini et al., 1991;

383 Carvalho *et al.*, 2007; Martinelli *et al.*, 2012) and the latest Cretaceous (*Peirosaurus*,

384 *Uberabasuchus*; Price 1955; Carvalho *et al.*, 2004). The latest Cretaceous diversity of

the group is also present in Madagascar, contributing to their higher diversity at the end-

386 Cretaceous in comparison with their initial peak of diversity in the Early Cretaceous

387 (Fig. 5.2).

388 Advanced notosuchians (the group to which *Notosuchus* belongs) are 389 exclusively known from South America and are virtually unknown prior to the 390 Turonian–Santonian. The only pre-Turonian record is *Coringasuchus anisodontis* from 391 the Cenomanian of Brazil (Kellner et al., 2009), a taxon known from fragmentary 392 remains and of uncertain affinities within the clade of advanced notosuchians (Pol et al., 393 2014). This group has a remarkable diversity during the Turonian–Santonian in South 394 America (over half of the notosuchian taxic diversity known from this period of time; 395 Fig. 5.2). This diversity is dominated by the numerous species of sphagesaurids known 396 from the Adamantina Formation of Brazil (Pol et al., 2014). After their diversity acme, 397 advanced notosuchians are rare and the only known advanced notosuchian from the 398 latest Cretaceous is Labidiosuchus amicum (Kellner et al., 2011b), a taxon known from 399 partial dentary remains from the Marilia Formation (Campanian-Maastrichtian) of 400 Brazil. The precise age of this diversity peak is slightly uncertain as there are authors 401 that propose the Adamantina Formation is younger in age than Turonian-Santonian (see 402 below) and because two other advanced notosuchians have been found in units with 403 poorly constrained ages (e.g., Late Cretaceous) of Bolivia (Yacarerani; Novas et al., 404 2009) and Argentina (Notosuchus n. sp.; Fiorelli et al., 2014). Therefore the diversity 405 peak of this group interpreted here as the Turonian–Santonian acme could be slightly

406 younger (toward the Campanian). As all other groups, advanced notosuchians become407 completely extinct by the end of the Cretaceous.

408 Baurusuchids are also known exclusively from South America and they are 409 completely absent from the fossil record prior to the Turonian–Santonian (Fig. 5.2). 410 Their sudden appearance occurs with high diversity levels, accounting for 411 approximately 30% of the taxic diversity known from this period of time. As in the case 412 of advanced notosuchians, this diversity is also dominated by the records from the 413 Adamantina of Brazil (Baurusuchus spp., Campinasuchus, Pissarachampsa, 414 Gondwanasuchus), with the addition some records from the Santonian (sensu Garrido, 415 2010) Bajo de la Carpa Formation of Patagonia (Fig. 5.2). After their diversity peak, 416 there are no definitive baurusuchids known in the fossil record. The two possible 417 exceptions are Pehuenchesuchus enderi from Patagonia (Campanian sensu Garrido, 418 2010) and Pabwehshi pakistanensis from Pakistan (Maastrichtian; Wilson et al., 2001), 419 but the former has uncertain affinities within Sebecosuchia (Pol et al., 2014) and the 420 latter has been retrieved as a peirosaurid by some phylogenetic analyses (e.g., Larsson 421 and Sues, 2007). Despite their possible survival until the end Cretaceous there are no 422 remains of this clade after the K/Pg boundary.

## 423 **DISCUSSION**

The diversity patterns shown above reveal the complex dynamics of notosuchian evolution during the Cretaceous, which highlights the components across space of two successive diversification pulses followed by two distinct extinction events that shaped the diversity curves of this clade. Despite the implemented phylogenetic and sampling corrections, these increases and drops in diversity are undoubtedly affected by the vagaries of the fossil record. Here we discuss these four events considering first the radiations and then the extinction events.

#### 431 Aptian Radiation

432 As shown above the first pulse of diversification in Notosuchia involves the 433 sudden appearance in the Aptian in the fossil record of three major clades, 434 Uruguaysuchidae, Peirosauridae (and allies), and basal lineages of Ziphosuchia (Fig. 435 5.2). These clades are the three most basal lineages of Notosuchia and their initial 436 diversification implies the existence of multiple ghost lineages during the Early 437 Cretaceous (Fig. 2), which led the proposal of this event as the Aptian radiation by Pol 438 et al. (2014). A problematic point related to this diversification event is the almost 439 exclusive absence of notosuchians during the earliest Cretaceous (Berriasian-440 Barremian), which hampers our understanding of the evolutionary dynamics of this 441 event (Pol et al., 2014). The absence of pre-Aptian notosuchians is most likely 442 influenced by the overall scarce fossil record of continental crocodyliforms during the 443 earliest Cretaceous, which contrasts with the denser sampling available for the Aptian-444 Albian (e.g., eight notosuchian bearing formations). Such situations may create an 445 artificial pattern of sudden radiation even when the diversification of these groups was 446 much more gradual and scattered through time. The only evidence suggesting this 447 indeed may be the case is Amargasuchus minor (Chiappe, 1988), known from a 448 fragmentary maxillary found at the Puesto Antigual Member of the La Amarga 449 Formation (Barremian). Amargasuchus was originally described as a member of 450 Trematochampsidae, a group of questioned validity but usually regarded as closely 451 related to or nested within Peirosauridae (Gasparini et al., 1991; Sertich and O'Connor, 452 2014). Furthremore, Gasparini et al. (1991) noted similarities between this fragmentary 453 form and the peirosaurid *Lomasuchus*. The putative existence of a peirosaurid, or even a 454 taxon closely related to this clade, in the Barremian would push the initial split of the 455 basal notosuchian lineages back to this stage. Furthermore, this would decouple the

456 diversification of peirosaurids from that of uruguaysuchids and basal ziphosuchians,

457 therefore questioning the existence of an Aptian radiation event. New information of the

458 earliest Cretaceous can easily influence current hypotheses on the initial diversification

459 of Notosuchia, even if they are based on fragmentary material such as the case of the

460 possible notosuchian *Amargasuchus*.

# 461 Turonian–Santonian Radiation

462 The major peak of notosuchian diversity is interpreted here as representing the 463 Turonian-Santonian radiation (Fig. 4), and the diversity curves split by geography and 464 subclades show this is largely formed by two South American groups: advanced 465 notosuchians and baurusuchids (Fig. 5). The well-sampled pre-Turonian units of 466 different regions of Gondwana, in which these two groups are absent, provide support 467 for interpreting this peak of diversity as a true radiation event during the Late 468 Cretaceous (Turner and Sertich, 2010; Pol et al., 2014). As noted by several authors 469 (e.g., Montefeltro et al., 2011; Pol et al., 2014; Martinelli and Teixeira, 2015), there is 470 some uncertainty related to the age of the Adamantina Formation, which provides over 471 80% of the known taxic diversity of advanced notosuchians and baurusuchids. Such 472 uncertainty may imply this radiation event occurred in South America during the 473 Santonian–Campanian rather than during the Turonian–Santonian. Despite these 474 uncertainties, it seems clear the existence of a remarkable radiation of baurusuchids and 475 advanced notosuchians (especially sphagesaurids) that shaped dynamics of the 476 terrestrial ecosystems in the middle Late Cretaceous in South America (e.g., Godoy et 477 al., 2014), especially in warm and dry (or seasonal) environments (Carvalho et al., 478 2010).

Whereas the existence of this radiation in South America is robustly supported,its endemic nature to South America cannot be adequately tested by the currently

481 available fossil record. Notosuchian remains from other regions during the Turonian-482 Santonian are extremely scarce, partly because there are no fossiliferous units of this 483 age in Africa, Madagascar, or other regions of Gondwana. Although not included in our 484 analysis, there are fragmentary remains of alleged notosuchian affinities from Europe 485 (Dalla Vechia and Cau, 2011; Rabi and Sebök, 2015) but these most likely do not 486 belong to the groups that radiated in South American during this time (*i.e.*, advanced 487 notosuchians and baurusuchids). The absence of relevant fossil and rock record 488 therefore precludes determining with certainty if advanced notosuchians or 489 baurusuchids were present in other regions of Gondwana during the middle Late 490 Cretaceous. Data from the latest Cretaceous (Campanian-Maastrichtian) of other 491 regions of Gondwana could provide hints regarding this point, but the available 492 information is inconclusive. On the one hand, the absence of advanced notosuchians or 493 baurusuchids in the diverse crocodyliform fauna of the latest Cretaceous Maevarano 494 Formation of Madagascar (Krause et al., 2006) suggests the South American endemism 495 of these two groups may be real, given that this unit records survival lineages of all 496 other notosuchian clades (e.g., uruguaysuchids, peirosaurids, basal ziphosuchians). On 497 the other hand, if the possible baurusuchid affinities of Pabwehshi from the 498 Maastrichtian of Pakistan are corroborated by more complete remains (e.g., preserving 499 the choanal region), it would significantly expand the geographic scope of this clade 500 across Gondwana. New data and crocodyliform remains from the middle Late 501 Cretaceous of Africa, Madagascar, and other regions of Gondwana will be critical for 502 adequately testing the South American endemism of the major radiation in the history 503 of Notosuchia.

504 Extinction patterns in Notosuchia

505 *Campanian event.* The first marked drop in diversity in the history of Notosuchia is 506 recorded at the beginning of the Campanian (Figs. 4–5) and is caused by the complete 507 extinction of baurusuchids and the drastic decrease in diversity of advanced 508 notosuchians in South America (Fig. 5). This decrease, at least in South America, seems 509 to be a real extinction event given that other groups of crocodyliforms are recorded in 510 the same geographic regions (e.g., southern Brazil, northwestern Patagonia). The above-511 mentioned uncertainty regarding the age of the Adamantina Formation and its impact on 512 the timing of the middle Late Cretaceous diversity peak also affects the timing of this 513 extinction event. If this formation is indeed younger than Turonian-Santonian as 514 proposed by some researchers (Gobbo-Rodrigues et al., 1999; Fernandes and Coimbra, 515 2000), this extinction event would have occurred later than the Campanian but before 516 the end of the Cretaceous Period. From a stratigraphic perspective, and given the 517 dominance of the Bauru Group in the notosuchian diversity curves, the extinction event 518 is nonetheless placed at the transition between the Adamantina and the Marilia 519 formations. The extent of this extinction event in other regions of Gondwana is largely 520 unknown due to the above-mentioned lack of Turonian-Santonian crocodyliform 521 record.

522 K/Pg event. Although the major decrease of notosuchian diversity occurred well before 523 the end of the Cretaceous Period, the K/Pg mass extinction event certainly affected 524 notosuchians. The five groups of Cretaceous notosuchians analyzed here (with the 525 possible exception of Baurusuchidae) were still present during the latest Cretaceous 526 (Fig. 5.2) but became completely extinct by the end of the Mesozoic. The only 527 notosuchians that survive this mass extinction event are sebecids (and related forms 528 recorded in the Paleogene of Europe and Africa; Buffetaut, 1989; Ortega et al., 1996; 529 Ortega, 2004). This group has been phylogenetically allied either to baurusuchids

(forming the clade Sebecosuchia; see Pol and Powell, 2011) or to peirosaurids (forming
the clade Sebecia; see Larsson and Sues, 2007). Irrespective of their debated affinities,
this distinct clade of notosuchians has its closest relatives in the Cretaceous and
therefore must have been originated prior to the K/Pg boundary. This is the only
notosuchian lineage that survived the mass extinction event (Gasparini, 1972, 1996;
Buffetaut, 1982; Kellner *et al.*, 2014) and subsequently became abundant during the
Paleogene, especially in South America (Fig. 5.1–2).

#### 537 CONCLUSIONS

538 The analysis of notosuchian diversity patterns reveals four basic episodes that 539 shaped the evolutionary dynamics of this group, involving two successive 540 diversification events and two consecutive extinction events that wiped out the 541 outstanding diversity of the group by the latest Cretaceous. The currently available 542 fossil record indicates the first diversification pulse (Aptian radiation) was 543 geographically widespread across Gondwana (and possibly other regions of the globe) 544 and involved the basal lineages of Notosuchia (Uruguaysuchidae, Peirosauridae and 545 allies, and basal forms of Ziphosuchia). However, the remarkably scarce record of pre-546 Aptian crocodyliforms suggests this may have not been a simultaneous event. In 547 contrast, the second pulse of diversification during the middle Late Cretaceous (possibly 548 Turonian–Santonian) is likely a true and outstanding radiation event (at least in South 549 America) of the highly diverse advanced notosuchians and baurusuchids. 550 Two consecutive but distinct extinction events are inferred to occur during the 551 latest Cretaceous (Campanian-Maastrichtian). The former of which records the almost 552 complete disappearance of advanced notosuchians and baurusuchids, leaving a 553 depauperated taxic diversity of notosuchians during the latest Cretaceous that were 554 subsequently affected by the end-Cretaceous mass extinction.

555 Different regions of Gondwana have provided critical evidence for 556 understanding the evolutionary pattern of Notosuchia at different periods of time during 557 the Cretaceous. South America dominates the notosuchian fossil record and basically 558 shapes the diversity curve during the entire Late Cretaceous. Most of what we currently 559 know about notosuchian evolution and diversity would be just impossible to infer without the contribution of the South American fossil record. The African fossil record 560 561 has been pivotal for understanding the initial diversification of the basal lineages of 562 Notosuchia and shapes the diversity curves during the "middle" Cretaceous (Aptian-563 Cenomanian). Finally, the latest Cretaceous record of Madagascar is relevant as it 564 shows the survival of lineages (*i.e.*, uruguaysuchids and basal ziphosuchians) that 565 became extinct in other regions of Gondwana earlier during the Cretaceous.

566 ACKNOWLEDGMENTS

567 This contribution aims to provide a synthesis of current knowledge of a group 568 whose modern understanding was founded by Zulma over 40 years ago, when she was a 569 young researcher working in her doctoral dissertation. Her influence in our field is 570 undeniable but Zulma has been even more important in terms of supporting, helping and 571 fostering the early careers of numerous researchers in Argentina. We therefore would 572 like to thank Zulma for her influence and help to us (and a large number of other 573 people) during the last years. We also would like to thank M. S. Fernández and Y. 574 Herrera for inviting us to contribute a chapter of this special issue, and for assembling 575 this issue dedicated to Zulma. Funds for this project were provided by the Agencia 576 Nacional de Promoción Científica y Técnica (PICT 2010-0756, 2013-0132, 2013-2725). 577 This is JML's R-166 contribution to the Instituto de Estudios Andinos Don Pablo 578 Groeber.

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883	
884	Figure captions
885	Figure 1. Skull reconstruction of four groups of Cretaceous notosuchians. 1,
886	Uruguaysuchidae (Araripesuchus gomesii); 2, Peirosauridae (Hamadasuchus
887	reboulii); 3, Notosuchidae (Notosuchus terrestris); 4, Baurusuchidae (Baurusuchus
888	salgadoensis). Modified from Pol and Larsson (2007).
889	
890	Figure 2. Number of notosuchian species discovered through time (blue curve). The pie

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- charts show the relative geographical distribution of the known diversity in 1974, 1994,
- and 2014, classified for major landmasses. References: blue: South America, red:
- 893 Africa, green: Madagascar; purple: other regions (China, Europe, Pakistan).
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Figure 3. Calibrated phylogeny of Notosuchia based on Pol *et al.* (2014) and Leardi *et al.* (2015). P\* indicates the clade formed by Peirosauridae and allies (*i.e.*,
Mahajangasuchidae).

898

**Figure 4.** Diversity curves of Notosuchia through time. **1**, Taxic diversity (dark grey)

900 and phylogenetically corrected diversity counting ghost lineages (light grey), with

901 vertical axis representing total number of species (taxic) or species plus ghost lineages

902 (phylogenetic correction) per period of time. 2, Taxic (dark brown) and

903 phylogenetically corrected (light brown) diversity corrected by sampling of fossiliferous

904 units, with vertical axis representing the taxic or phylogenetic diversity divided by the

905 number of notosuchian bearing formations per unit of time.

906

907 Figure 5. Taxic diversity of Notosuchia split by 1, geographical regions and 2, different

908 notosuchian groups across time. For each period of time the different contributions of

909 each region or clade are on top of each other so that the sum equals the taxic diversity

shown in Figure 4.1.