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Title: Niche overlap and competition potential among tigers (Panthera tigris), sabertoothed cats (Homotherium ultimum, Hemimachairodus zwierzyckii) and Merriam's Dog (Megacyon merriami) in the Pleistocene of Java

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Keywords: carnivores; body mass; ecological character displacement; prey mass spectrum; Sangiran; Southeast Asia

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Abstract: On Java during the Pleistocene, tigers of more than 300 kg occurred, but these are restricted to a single Late Pleistocene faunal unit, while Early and Middle Pleistocene tigers possessed body masses comparable to those of historic Javanese and extant Sumatran tigers. The aim of this study is to test if competition for prey with other hypercarnivorous taxa such as sabertoothed cats and the large Merriam's Dog was the driver for the increase in body mass of tigers. We calculated body masses and prey mass spectrum for tigers and potential competitors using linear regressions. Niche overlap was then estimated based on the prey mass spectrum after which niche overlaps were used as indicators for competition potentials. Reconstructed body mass for Homotherium ultimum, Hemimachairodus zwierzyckii, Megacyon merriami are 154 kg (comparable to Homotherium from Untermassfeld), 130 kg and 52 kg, respectively. The niche overlap between tigers and Merriam's Dog is highest (100%) while it is comparatively low (60 %) between tigers and Homotherium ultimum. In order to reduce competition, tigers seem to have increased body mass to avoid competition especially with Merriam's Dog whereas Merriam's Dog on its turn seems to have decreased body mass to avoid competition with tigers. The sabertoothed cats on the other hand seem to have been unable to adapt and went extinct.

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21 Abstract

22 On Java during the Pleistocene, tigers of more than 300 kg occurred, but 23 these are restricted to a single Late Pleistocene faunal unit, while Early and 24 Middle Pleistocene tigers possessed body masses comparable to those of 25 historic Javanese and extant Sumatran tigers. The aim of this study is to test if 26 competition for prey with other hypercarnivorous taxa such as sabertoothed 27 cats and the large Merriam's Dog was the driver for the increase in body mass 28 of tigers. We calculated body masses and prey mass spectrum for tigers and 29 potential competitors using linear regressions. Niche overlap was then 30 estimated based on the prey mass spectrum after which niche overlaps were 31 used as indicators for competition potentials. Reconstructed body mass for 32 Homotherium ultimum, Hemimachairodus zwierzyckii, Megacyon merriami are 33 154 kg (comparable to *Homotherium* from Untermassfeld), 130 kg and 52 kg, 34 respectively. The niche overlap between tigers and Merriam's Dog is highest 35 (100%) while it is comparatively low (60%) between tigers and Homotherium 36 *ultimum.* In order to reduce competition, tigers seem to have increased body 37 mass to avoid competition especially with Merriam's Dog whereas Merriam's 38 Dog on its turn seems to have decreased body mass to avoid competition with 39 tigers. The sabertoothed cats on the other hand seem to have been unable to 40 adapt and went extinct.

41 1 Introduction

42 Body masses of tiger subspecies vary in accordance with latitude and 43 geographic distribution (Mazák, 1981). In Southeast Asia body masses of 44 tigers, for example, range from 80 to 120 kg while body masses of Siberian 45 tigers can reach up to 300 kg (Mazák, 1981). This variation in body masses is 46 regularly explained with Bergmann's rule according to which closely related 47 mammalian taxa tend to be larger in colder environments than in warmer ones 48 (Bergmann, 1847; Seidensticker et al., 1999; Meiri et al., 2007). 49 In the Pleistocene, however, tigers with body masses in excess of 300 kg also 50 occurred in lower latitudes, particularly in Java (Hertler and Volmer, 2008). 51 Such high body masses cannot be understood as an adaptation to cold climate in the sense of Bergmann. Moreover, the large tiger individuals are 52 53 restricted to a single Late Pleistocene faunal unit Ngandong (Yokoyama et al., 54 2008), while Early and Middle Pleistocene tigers of Java possess body 55 masses comparable to historic Javanese and extant Sumatran tigers (Hertler 56 and Volmer, 2008). 57 Competition with other hypercarnivorous taxa may provide an alternative 58 explanation for the observed shifts in body mass. The prey mass spectrum of 59 carnivores strongly correlates with body mass (Carbone et al., 1999; Hemmer, 60 2004) and shifts in body mass lead to shifts in prey mass spectra (Sinclair et 61 al., 2003). Extensive overlap in the prey mass spectra among any pair of 62 coexisting carnivores can therefore be reduced by increase or decrease in 63 body mass. 64 Body mass increases observed for tigers in the Late Pleistocene Ngandong

65 faunal stage may therefore have been induced by a strong niche overlap with

66 the short-faced hyena (Pachycrocuta brevirostris) during the preceding faunal 67 unit (Hertler and Volmer, 2008). Consequently, this would have led to a reduction of competition. Niche overlap between tigers and other competitors 68 69 like Trinil Dog (Mececyon trinilensis, or Cuon trinilensis in Louys, 2014), 70 leopard (*Panthera pardus*) and the Asiatic wild dog (*Cuon alpinus*) are much 71 lower on the ground of their much lower body masses (Hertler and Volmer, 72 2008). These taxa are therefore not considered as ecologically significant 73 competitors for tigers in the Ngandong faunal unit. 74 Hertler and Volmer (2008) also mentioned additional large carnivore taxa that 75 co-occurred with tigers during the Pleistocene of Java, namely the 76 machairodonts Homotherium ultimum and Hemimachairodus zwierzyckii, as 77 well as a very large canid, Megacyon merriami (von Koenigswald, 1940; 78 Schütt, 1973). This raises the question whether competition between tigers 79 and the machairodonts and canids may have had an additional impact on tiger 80 body masses. 81 The succession of mammalian faunas at Sangiran illustrates a biostratigraphic 82 sequence (Fig. 1), which calibrates other Javanese fossil localities (Fig. 2) in 83 Central and East Java (Watanabe and Kadar, 1985). Von Koenigswald 84 collected a large number of mammalian fossils from Sangiran, including 85 specimens attributed to the machairodonts and Merriam's Dog (von 86 Koenigswald, 1940). However, the deposits at the Sangiran dome cover a 87 lithostratigraphic sequence extending possibly over a million years (Watanabe 88 and Kadar, 1985; Sémah et al., 2003). The lack of crucial stratigraphic 89 background information in Von Koenigswald's collection hampers the 90 assessment with whom Homotherium ultimum, Hemimachairodus and

91 Megacyon competed. Beside fossils collected by Von Koenigswald in 92 Sangiran, fossils from Teguan and Jeruk are found by Dubois and cannot be 93 attributed to one of the faunal levels (van den Bergh, 2001). Among these 94 fossils are also an ulna from a tiger found at Teguan and a maxilla found in 95 Jeruk (Brongersma, 1935). Reconstruction of body masses may help to 96 assign fossils from Teguan and Jeruk to a faunal level. Nevertheless, it is evident that the machairodonts as well as Merriam's Dog 97 98 were collected from Early and/or Middle Pleistocene deposits. Von 99 Koenigswald (1934, 1974) further informs that Megacyon merriami was 100 collected from the black clays (von Koenigswald 1940). The black clays at Sangiran encompass both the Satir and Ci Saat faunal unit (Watanabe and 101

102 Kadar, 1985). The Satir faunal unit represents an imbalanced island fauna

103 without carnivores (Sondaar, 1984), which implies that Merriam's Dog

104 occurred in the Ci Saat faunal unit. The machairodonts should be attributed to

105 either the Ci Saat faunal unit or the Kedung Brubus faunal unit. For

106 stratigraphic reasons, the Trinil H.K. faunal unit, which is bracketed by Ci Saat

107 and Kedung Brubus faunal units, cannot be excluded either for the

108 machairodonts. With respect to biostratigraphy, tigers are likely part of all

109 three faunal units (Fig. 2). Tigers must thus have competed with the

110 sabertoothed cats and Merriam's Dog.

111 Here, we will reconstruct body masses and prey mass spectra for all large

112 carnivores from Sangiran in order to address potential niche overlaps between

113 tigers (Panthera tigris), sabertoothed cats (Homotherium ultimum,

114 Hemimachairodus zwierzyckii) and Merriam's Dog (Megacyon merriami). Our

115 hypothesis is that extensive niche overlap among these taxa indicates strong

competition which subsequently resulted in body mass shifts observed in laterfaunal units.

118 2 Material and Methods

Body masses were reconstructed for canids and felids from Sangiran (based

120 on regressions; details in Appendix A). Based on these body masses, we infer

121 prey mass spectra (PMS) using a second set of regressions. Finally, niche

122 overlaps (NO) between the taxa at Sangiran are reconstructed. Broad niche

123 overlap is used as a proxy for high competition potentials, because prey of the

124 same size is consumed by both competitors.

125

126 The fossil samples are stored in the Von Koenigswald collection

127 (Senckenberg Research Institute, Frankfurt am Main, Germany), the Dubois

128 Collection (Naturalis Biodiversity Center, Leiden, the Netherlands) and the

129 Museum of Geology (Bandung, Java) (Error! Reference source not

130 **found.**)Error! Reference source not found..

131 The tiger samples for reconstructing body mass come from the following sites:

132 Dubois Collection: Trinil (n=13), Pitu (n=1), Jeruk (n=1), Kebon Duren (n= 1),

133 Kedung Brubus (n=2), Teguan (n=1), von Koenigswald Collection: Sangiran

134 (n=14) and the Museum of Geology: Bumiayu (n=2), Ngasinan (n=1),

135 Ngandong (n=7), and Watualang (n=2). The sites of Ngandong, Ngasinan and

136 Watualang belong to the Ngandong Faunal stage (Weidenreich, 1951;

137 Sondaar, 1984; Hertler and Volmer, 2008), whereas all other samples are

138 classified as pre-Ngandong-samples. The samples from Pitu and Trinil are

139 classified as Trinil H.K. sample and Kedung Brubus and Kebon Duren as

140 Kedung Brubus sample. All samples are shown in Table 2 and their

141 stratigraphic background in Table 1. The 40 tiger fossils from Sangiran

142 (Hemmer, 1971) originate from Early and/or Middle Pleistocene deposits.

143 Because of the absence of stratigraphic background we excluded specimens

144 from the Sangiran dome from our previous studies (Hertler and Volmer, 2008).

145 Nonetheless, all specimens were retrieved from deposits at Sangiran, which

implies that they are geographically congruent and are thus included in thisstudy.

148 The samples of the *Homotherium ultimum* (n=1), *Hemimachairodus* 

149 *zwierzyckii* (n=1) and *Megacyon merriami* (n=3) are stored in the Von

150 Koenigswald-Collection and originate from the Sangiran Dome. The

151 machairodonts and *Megacyon merriami* are here attributed to the Ci Saat

152 faunal stage (see section 1).

153

154 2.1 Body mass and prey mass spectrum reconstructions

155 For reconstruction of body masses on the basis of long bones (humerus,

156 femur, tibia, ulna) the regressions of Christiansen and Harris (2005) are used.

157 Since for teeth only regressions based on M1 length are published, we

158 calculated regressions for other tooth positions based on samples of extant

159 felids and canids. We also calculated a tiger specific regression based on the

160 correlation between skull length and body mass in extant tiger subspecies.

161 The samples and the procedure of calculating these regressions (Error!

162 **Reference source not found.**) can be found in the supplementary material of

163 this paper (Appendix A Supplementary data).

164 Body mass is reconstructed for each fossil. For skulls, mandibles and teeth,

165 the computed regressions as given in Error! Reference source not

166 found.Error! Reference source not found.Table 3 are used. For mandibles and

167 skulls, body mass is estimated for each usable parameter after which the

168 means of all estimated body masses was calculated for the respective

169 specimen.

170 For each sample the mean body mass was calculated. For Javanese tigers,

the mean body mass of the pre-Ngandong and Ngandong samples were

172 compared by a t-test. In addition, the means of the samples from Trinil,

173 Kedung Brubus, Sangiran and Ngandong were compared with ANOVA.

174 Each sample/taxon is then characterized by its minimum (BM<sub>min</sub>), mean

175 ( $BM_{mean}$ ) and maximum body mass ( $BM_{max}$ ). We applied the mean percentage

176 prediction error (%PE) to calculate a range of reconstructed body masses. For

each sample the minimum and the maximum body mass including errors were

178 determined (BM<sub>min%PE</sub> und BM<sub>max%PE</sub>).

179 All taxa included in this study have a hypercarnivorous diet, according to the

180 definition of Van Valkenburgh (1988), who defined a hypercarnivous diet as a

181 diet of which at least 70% of consumed biomass comes from self-hunted

182 vertebrate prey (van Valkenburgh, 1988, 2007; Wesley-Hunt, 2005).

183 Felids are characterized by hypercarnivorous dentitions; especially

184 machairodonts show a kind of "carnassialisation" of their premolars (Thenius,

185 1989; Ewer, 1998). Accordingly, the tiger and the maicharodonts are treated

186 as hypercarnivores in this study.

187 In case of canids, the diet varies from hypocarnivorous (e.g. *Chrysocyon*) to

188 hypercarnivorous (e.g. *Lycaon pictus*) and this is reflected in both tooth and

skull morphology (Ewer, 1998). Dental morphology can be used to reconstruct
the diet of fossil taxa (Ewer, 1998). Lyras et al. (2010) consider Merriam's Dog
as hypercarnivorous, because its dentition is most comparable to the
hypercarnivorous morphology of the molars of *Xenocyon* and *Lycaon pictus*.
Also Schütt (1973) mentioned the similarity of *Megacyon* to *Xenocyon* and
that both likely belong to the same genus.

195 Hunting strategies may also differ and can have a significant impact on the

196 prey mass spectra in canids (Nudds, 1978). Most hypercarnivorous canids

197 hunt in packs. Due to energetic constraints, carnivores with body masses of

198 21 kg or more hunt on prey larger than themselves (Carbone et al., 1999;

Bogusch, 2002), which is achieved by canids by hunting in a group (van

200 Valkenburgh et al., 2003). Therefore, we compare *Megacyon* here with the

201 pack-hunting hypercarnivorous canid *Cuon alpinus*.

202 Prey mass spectrum is required in order to calculate the potential niche

203 overlap. The prey mass spectrum of carnivores is often demonstrated by body

204 mass classes following logarithmic steps (Hemmer, 2004; Hertler and Volmer,

205 2008). The total hunted prey biomass is distributed over these body mass

206 classes. Some classes contribute relatively more to the total hunted prey

207 biomass than others (Volmer, 2013). These classes are the focus classes of

the top predators and it is assumed that the predator mainly subsists on prey

209 of these classes. The focus classes can be reconstructed using regressions

terms to calculate the mean prey mass (Hemmer, 2004; Hertler and Volmer,

211 2008).

212	A regression term is known only for felids (Hemmer, 2004), whereas such a
213	regression is missing for canids. But also in case of the regression term for
214	felids, it remains unclear on which sample it has been computed.
215	In order to reconstruct the prey mass spectra of Felidae and Canidae, we
216	computed new regressions based on the correlation between body mass and
217	prey mass of predators using data and samples of field studies of extant
218	members of these families. These regressions were calculated for felids and
219	canids separately, because both follow different hunting techniques. In case of
220	felids we included only large cats (> 10 kg) in our sample, because small cats
221	use different hunting styles (Schaller, 1972; Leyhausen, 1979; Ewer, 1998)
222	focusing on prey smaller than themselves (Carbone et al., 1999).
223	The mean prey mass of each taxon is calculated from prey counts of several studios listed in
224 225	Table 4. The advantage of using more than one study from one study area to
226	calculate the mean body mass of a species, is that the influence of other
227	factors - e.g. prey offer, small sample size of kills/scats - which affect the
228	prey mass spectrum can be reduced. The body masses of the comparative
229	predators are taken from the literature (Jerdan, 1984; Kingdon, 1997;
230	Eisenberg and Redford, 1999; Smith and Xie, 2008).

- 231 The focus classes were calculated from the calculated regression terms (
- Table 5) and the reconstructed body masses from
- 233 Table 6. The mean prey mass was calculated by the regression term and determines
- the focus class. Prey masses were calculated for each species using the minimum
- 235 (BMmin), average (BMmean), maximum (BMmax) body mass. To account for the
- statistical error of body mass reconstructions the prey mass was also calculated for
- 237 BM<sub>min%PE</sub> and BM<sub>max%PE</sub> and prey mass classes will be assigned. These are called
- 238  $FC_{Min}$ ,  $FC_{Mean}$ ,  $FC_{Max}$  and  $FC_{Min\%PE}$ ,  $FC_{Max\%PE}$ .
- 239 Since the regression terms for calculation of prey masses also yield statistical
- 240 errors, the reconstructed focus classes were compared to prey mass spectra
- of comparable extant family members with comparable size and a
- 242 hypercarnivorous diet.
- 243
- 244 2.2 Calculation of niche overlap (NO)
- Niche overlap was calculated as the percentage of overlap following
- 246 Renkonen and Schoener (Renkonen, 1938; Schoener, 1970; Krebs, 1999),
- which is a measure of the actual area of overlap of the resource utilization
- curves of two species (Krebs, 1999).

249 Its equation is:

- 250  $p_{jk} = [L_j = \sum_{i=1}^{n} (p_{ij}, p_{ik})] \times 100$ , where
- 251  $p_{ik}$  = percentage overlap between species *j* and species *k*
- 252  $p_{ij}$  = proportion resource *i* is of the total resources used by species *j*
- 253  $p_{ik}$  = proportion resource *i* is of the total resources used by species *k*
- n = total number of resource states

- 255 In this study the resources were reflected by the prey mass classes according
- to Hertler and Volmer (2008). The total of focus classes for a sample is

assumed to represent 100% and each focus class contributes equally. For

- example, if the reconstructed focus classes of predator A are 3a, 3b, 3c and
- 4a, and those of predator B are 3c, 4a, 4c, 5a and 5b, it follows that predator
- A experiences a niche overlap of 50% with predator B, but predator B
- 261 experiences an niche overlap of only 20% by predator A.
- 262 The higher the niche overlap, the higher is the potential competition between
- the competing taxa.
- 264 3 Results
- 265 3.1 Reconstructed body masses
- 266 Reconstructed body masses are shown in
- Table 6 and Figure 3. Sangiran tigers have a mean body mass of 121 kg and
- 268 differ not significantly from other pre-Ngandong tigers. The largest Sangiran
- tigers virtually reach the mean body mass of Ngandong tigers (182 and 184
- kg, respectively).
- 271 The means of the single pre-Ngandong samples (Bumiayu, Trinil H.K.,
- 272 Kedung Brubus and Sangiran sample) vary between 87 and 122 kg.
- 273 The mean body mass for the Ngandong sample is 184 kg and for the pre-
- Ngandong sample 114 kg. The Ngandong tiger is significantly larger than the
- pre-Ngandong tiger regarding its body mass (ANOVA p=0,0073, Table 8). The
- 276 reconstructed body mass of *Homotherium ultimum*, *Hemimachairodus*
- zwierzyckii and Megacyon merriami is 154 kg, 130 kg and 52 kg respectively.

- 279 3.2 Prey Mass Spectra and niche overlaps
- 280 The regressions for mean prey mass reconstruction show correlations
- between mean body mass and prey mass (
- Table 5). The prey mass spectra are shown in
- Table 7. pre-Ngandong tigers and Sangiran tigers focus on the same prey
- classes (3a to 4b). The prey mass spectrum of Ngandong tigers is shifted and
- focusses on classes 3b to 5a. *Megacyon merriami* focuses on classes 3b to
- 4b which coincide with those of Sangiran tigers.
- 287 Homotherium ultimum and Hemimachairodus zwierzyckii focus on classes 3c
- to 4c and 3b to 4b, respectively.
- 289 Prey mass spectra of Merriam's Dog overlap strongest with Sangiran and pre-
- 290 Ngandong tigers (NO = 100%) and vice versa (Figs 4 and 5).
- 291 Hemimachairodus also overlaps with the tiger by 100%, but Homotherium
- 292 only by 60%. Therefore, the highest competition potential is observed
- between Sangiran tigers and pack-hunting Merriam's Dog.
- 294 Niche overlaps between tigers from Ci Saat, Trinil H.K. or Kedung Brubus and
- the three other carnivores from Sangiran show slightly different results:
- 296 Merriam's Dog and *Hemimachairodus* both overlap all tigers by 100%.
- 297 Homotherium ultimum overlaps the tiger from Bumiayu by 66–67% and both,
- the Trinil H.K. and Kedung Brubus tigers by 75%.
- 299 Niche overlap experienced by the competitors show similar results. *Megacyon*
- 300 *merriami* is overlapped by the Sangiran tiger by 100% (Fig. 5). The
- 301 Machairodonts experience smaller niche overlaps (60–80%) by the Sangiran
- 302 tiger. Niche overlaps with the Ngandong tiger are smaller for *Megacyon*
- 303 merriami and Hemimachairodus zwierzyckii, but larger for Homotherium
- 304 ultimum.

# 305 4 Discussion

# 306 4.1 Reconstructed body masses

307	All regressions for body mass reconstruction show a significant correlation
308	between tooth/skull length and body mass. The strongest correlations are
309	found for the carnassials (lower M1 and upper P4) in both felids and canids.
310	In felids, the upper P2 gives strong statistical errors ( $r^2 = 0.314$ , %SEE = 177,
311	%PE = 76%). The reason for this remains unclear, but this premolar is
312	reduced in felidsin some species it is even completely lackingand shows
313	generally a high variability in its morphology (Thenius, 1989; Ewer 1998).
314	Tooth positions with a high variability usually show low correlation with body
315	mass (van Valkenburgh, 1990). Thus we excluded the upper second
316	premolars from the body mass reconstructions in this study and cannot
317	recommend using this tooth position in other studies.
318	Regressions based on dental elements have on average higher statistical
319	errors in comparison to the regressions for long bone elements developed by
320	Anyonge (1993) and Christiansen and Harris (2005). Christiansen and
321	Harris's regressions (2005) show %SEEs of 16–55 and Anyonge regressions
322	show %SEEs of 24–39 for felids and 23–41 for canids (1993). Our %SEE for
323	felids range from 14–50 % with the exception of the upper P2. The
324	regressions based on skull length of tigers, lower M1 and upper P4 have
325	statistical errors that fall into the range of error of limb bone regressions.
326	Therefore, we consider these two tooth positions (the carnassials) as well as
327	skull length as reliable as limb bones for body mass reconstruction.

328 %SEE of the canid regressions for lower P3, lower P4 and lower M1 also fall 329 within the range of the limb bone regressions by Anyonge (1993), which makes them in our view as reliable as the limb bone measurements. 330 331 If tooth morphology differs strongly from that in the sample used for the 332 calculation of the regressions, the reconstructed body mass may vary widely 333 from its actual body mass, since teeth are no weight bearing structures 334 (Fortelius, 1990). This may apply to the reconstructed body masses of 335 Homotherium and Hemimachairodus. The lower P3 of Homotherium ultimum 336 is strongly reduced and we therefore excluded this position for the 337 reconstruction of its body mass. Application of the regression equation to the 338 lower P3 of *H. ultimum* would result in a body mass of only 24 kg. Obviously, 339 this value underestimates body mass, since other body mass reconstructions 340 of Homotherium crenatidens from Untermassfeld (Early Pleistocene, 341 Germany) range between 180 kg and 300 kg and those of *Homotherium* 342 serum range between 134 kg and 236 kg (Anyonge, 1993; Hemmer, 2001). The 218 kg for *Homotherium ultimum* from Java as the means of regressions 343 344 based on the carnassials thus confirms the estimations for the other two 345 species of Homotherium from Untermassfeld. 346 Estimated body masses for Merriam's Dog vary around 52 kg which is 347 comparable to the size of extant gray wolves (*Canis lupus*). The reconstructed 348 masses based on measurements of the upper and lower first molars do not 349 differ much (49–55 kg) and thus both of them have been applied for the 350 reconstruction of prey mass spectra.

The body masses of tigers do not differ significantly among the pre-Ngandong
 samples. Variation in body mass is low, although some fossils from Sangiran

353 fall in a range above the mean body mass of Ngandong tigers. This illustrates 354 that single large individuals existed already in the Middle Pleistocene, but the majority of individuals was smaller than those of the Late Pleistocene. 355 356 The wide range of body masses in Ngandong tigers likely reflects sexual 357 dimorphism of these tigers. Females of all subspecies do not differ strongly in 358 body size in contrast to males (Seidensticker et al., 1999). This sexual 359 dimorphism increases with size, for example males are 185% larger than 360 females in P. t. tigris and 206% larger in P. t. altaica (Mazák, 1981). 361 Our study confirms the increase in tigers' body mass from the Middle to the 362 Late Pleistocene. The t-test comparing the mean reconstructed body mass of 363 pre-Ngandong and Ngandong samples indicates a significant difference at the 364 1%-level (p= 0.0073).

365 Two tiger fossils from the sites Teguan and Jeruk could not be attributed to a certain faunal stage in earlier studies (van den Bergh et al., 2001). The tiger 366 367 fossil of Teguan has a reconstructed body mass of 86 kg and is comparable 368 with body masses reconstructed from the pre-Ngandong sample. In addition, 369 the presence of *Pachycrocuta brevirostris* in Teguan suggests an attribution of 370 this site to the Kedung Brubus faunal stage (de Vos pers. comm.). The 371 reconstructed body mass does not exclude this, but cannot confirm the 372 assumption either.

On the other hand, the tiger fossil of Jeruk has a reconstructed body mass of 228 kg. This body mass only falls in the range of the Ngandong tiger sample, and thus suggests the attribution of Jeruk to the Ngandong faunal stage. The presence of *Panthera pardus* at this site (Brongersma, 1935; Hemmer and

Schütt, 1972) further confirms this, because *Panthera pardus* is known only
from the Ngandong faunal stage and onwards.

379

380 4.2 Prey mass spectra (PMS)

381 The regressions for reconstruction of the mean prey masses show high

382 statistical errors (%PE 68 and 95, %SEE 138 and 192). Since other studies

383 about correlations between body mass and prey mass do not provide any

information about their statistical errors (Carbone et al., 1999; Hemmer, 2004)

385 we cannot compare our regressions with similar ones.

In order to cope with this shortcoming and to reduce the errors, we applied a
 classification system. In addition, we compared the reconstructed focus

388 classes with results from comparable members of the respective families. This

389 method can also be used on itself for estimating the focus classes of fossil

390 carnivores. However, in some cases data for species or comparable family

391 members are unavailable. In addition, in case of fossil taxa we cannot be

392 certain about their hunting behavior. As we have shown here, Merriam's Dog

393 likely was a pack hunter.

394 In the case of extant cats, hunting strategy has no strong effect on prey mass 395 spectrum (Hemmer, 2004). The only pack hunting felids are lion and cheetah, 396 whose females or males respectively hunt in packs. Calculation of the prey 397 mass on the basis of regression F-1 would lead to an overestimation by 24% 398 for the lion (predicted: prey mass 295 kg, prey class 4b; observed: prey mass 399 238 kg, prey class: 4b) and for the cheetah to a underestimation of 46 % 400 (predicted: prey mass 20 kg, prey class: 3b; observed: prey mass 36 kg, prey 401 class: 3b). Since over- and underestimation occur in solitary taxa as well, we

402 cannot conclude that pack hunting on itself leads to an underestimation of
403 prey mass by regression F-1. In both cases, the prey class of the observed
404 and predicted prey mass is identical and illustrates that the regression is
405 applicable for reconstruction of prey mass spectra.

Whether machairodont taxa like *Homotherium* and *Hemimachairodus* were
pack hunters or solitary hunters will probably never be revealed. *Homotherium latidens* is generally considered as a pack hunter, because of its cursorial
adaptations and reduced claws (Antón et al., 2005). *Hemimachairodus* is only
known from the Pleistocene of Java and no postcranial material was found.
But even if it would have focused on prey larger than reconstructed, this would

412 only have further reduced its competition potential with the tigers.

413

414 4.3 Competition potential among Sangiran carnivores and its impacts 415 The Sangiran tiger and Pre-Ngangdong tiger have the same prey mass 416 spectra and thus also the same niche overlaps with the other three potential 417 competitors. The pre-Ngandong sample includes body masses from all Middle 418 Pleistocene sites in Java where the tiger occurred. Thus, there was no 419 geographic variation in body mass and prey spectrum. The pre-Ngandong 420 tiger focuses on classes 3a-4b according to prey from 10-200 kg. 421 From pre-Ngandong to Ngandong we can observe a shift to the classes 3b-422 5a (20–1000 kg). This shift in the prey mass spectrum is caused by the shift in 423 body masses of the Ngandong tiger. Thus, the increased body mass of the 424 tiger had an impact on its prey spectrum and thus consequences for 425 competition relations, independent of whether competition was the main 426 driver for the body mass increase or not. The prey mass spectrum of the

427 Ngandong tiger includes class 5a, which is not hunted by any other potential 428 competitor. This applies also to Pachycrocuta brevirostris, Cuon alpinus and Panthera pardus whose prey mass spectra were reconstructed in our former 429 430 study (Hertler & Volmer, 2008). Further, its prey mass spectrum includes class 4c which is hunted only by *Homotherium ultimum*. The Ngandong tiger thus 431 432 has the lowest niche overlap with *H. zwierzyckii* and Merriam's Dog. 433 It focuses on the same prey classes as *Megacyon merriami* and thus 434 experienced a niche overlap of 100% by *M. merriami*. The Ngandong tiger 435 would have experienced a smaller niche overlap by Merriam's dog (67%) than 436 the pre-Ngandong tiger. Thus, the increase in the body mass of the tiger 437 would have lowered the competition potential between both species. 438 Merriam's Dog also experienced a niche overlap of 100% by the tiger. As a 439 solitary hunter, Megacyon would have had no niche overlap with tigers at all. It 440 could have avoided niche overlap and competition with tigers by adopting a 441 solitary hunting style. More likely is the strategy proposed by Lyras et al. (2010), who suggest that Merriam's Dog eventually reduced its body size and 442 443 evolved into the jackal-sized *Mececyon trinilensis*. In this way, there is no 444 competition with tigers. An alternative scenario is that Megacyon could not 445 adapt, went extinct and was replaced by a second, much smaller 446 hypercarnivorous canid with no niche overlap with the tigers. 447 Both scenarios are supported by our data, because Merriam's Dog had the 448 same reconstructed focus classes as the tiger whereas that of *Mececyon* falls 449 well below this class. The derived *Mececyon* would not have suffered from 450 niche overlap with the other carnivores from Sangiran because the clouded

451 leopard is significantly larger and the Bengal cat, mustelids and viverrids452 significantly smaller.

453 Competition between tigers and Homotherium ultimum is not likely to have 454 had an impact on the extinction of the sabertoothed cats or shifts in body 455 masses of the tigers. Homotherium ultimum has the largest body mass and 456 thus focused on larger prey classes having a PMS of 3c-4c. The pre-Ngandong tiger got overlapped by Homotherium only by 60%. This is the 457 458 lowest niche overlap the pre-Ngandong tiger would experience if it would have 459 competed to Sangiran carnivores. In contrast, the Ngandong tiger would have 460 been overlapped by Homotherium ultimum by 83%. This is 23% more and 461 thus, the increase in the body mass of the tiger likely did not evolve in 462 response to competition with Homotherium ultimum. 463 Hemimachairodus zwierzyckii shows an overlap of 80% with Sangiran and pre-Ngandong tigers. This is as strong as the short-faced hyena and tigers in 464 465 Kedung Brubus (Hertler and Volmer, 2008). The observed increase in body 466 mass of the tiger in the Late Pleistocene would lead to a decrease in niche 467 overlap to 67%. Thus, competition between Hemimachairodus zwierzyckii 468 may have been the reason for body mass increase in the tiger to avoid 469 competition. 470 In summary, in the Ci Saat faunal level the coexistence of Merriam's Dog,

Hemimachairodus zwierzyckii and Panthera tigris implied a high degree of
competition which likely led to competition avoidance strategies. This is in
case of Hemimachairodus zwierzyckii its extinction, while Panthera tigris
increased its body mass and thus shifted its prey mass spectrum. Indeed,
Hemimachairodus zwierzyckii is already extinct in the following Trinil H.K.

476 faunal level. Merriam's Dog on the other hand was replaced by *Mececyon* 

477 *trinilensis* in the Trinil H.K. faunal unit. The extinction of *Homotherium ultimum* 

478 cannot be explained by competition between any of the other large carnivores.

479

480 Conclusion

481 Our results confirm that tigers increased their body mass between the Middle

and Late Pleistocene. While tigers from the sites of Teguan have body

483 masses comparable to those of pre-Ngandong tigers, the tiger from Jeruk falls

in the range of the large Ngandong tigers and suggests an attribution of this

485 site to the Ngandong faunal stage.

486 There was no strong niche overlap between tigers and *Homotherium ultimum* 

487 which makes it unlikely that competition with *Homotherium* forced tigers to488 increase their body masses.

489 Interestingly, Merriam's Dog shows complete niche overlap with the tiger and

490 this constituted a high competition potential. After the short-faced hyena and

491 *Hemimachairodus zwierzyckii*, Merriam's Dog had the highest competition

492 potential. The hypothesis that Merriam's Dog was replaced by the much

493 smaller Trinil Dog, which had a lower niche overlap with the tiger, is thus

494 supported by our data.

495 If competition forced the tiger to increase its body mass, the short-faced

496 hyena and/or Merriam's Dog followed by Hemimachairodus zwierzyckii are

the most parsimonious candidates with high competition potential for the tiger.

498

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Table 1 Faunal stages and the samples for body mass reconstruction and 690 691 calculation of niche overlaps. Shown are the carnivora taxa for each faunal stage after van den Bergh et al. (2001); Hertler and Volmer (2008) and Lyras 692 693 et al. (2010) and the attributed samples. For the felid samples from Sangiran, 694 the exact faunal stage remains unknown (see explanation in text). The Pre-695 Ngangdong sample is a summary of the Sangiran Sample, the attributed Tiger sample. The samples Teguan and Jeruk cannot be attributed to one of the 696 697 faunal stages. The fossils included in the samples are shown in Table 2. The 698 biostratigraphic sequence is based on Sondaar (1984), Leinders et al. (1985) 699 and includes later revisions by de Vos and the Long (2001) and van den 700 Bergh et al. (2001). 701 \* Swisher et al. (1996) suggests dates from 27–53 ka based on electron-spin 702 resonance and U-series dating of fossil bovid teeth. However, Westaway et al. 703 (2007) dated the subsequent Punung fauna to the Last interglacial (between 704 128 +/- 15 and 118 +/- 3 ka) and conclude that the Ngandong Fauna must be 705 older than the Last Interglacial. Yokoyama et al. (2008) dated the Homo 706 erecuts skulls from Ngandong to around 40 ka, with an upper limit of around

707	60 to 70 kg	by gamma ray	( apostromatry)
/0/	00 10 7 0 Ka	by gamma-ray	y specifornelly

Faunal Stages	Age	Carnivora taxa	Sangiran stratigraphy	Samples		
Ngandong	Late	Panthera tigris	sterile layers	Ngandong-		
	Pleistocene*			Sample		
		Panthera pardus		(Panthera		
				<i>tigris</i> n=10,)		
		Cuon alpinus		Sites:		
				Ngandong,		
				Ngasinan,		
				Watualang	-	
Kedung	0,7-0,8 ma	Panthera tigris	upper Bapang	Kedung		Pre-
Brubus				Brubus		Ngangdong
		Pachycrocuta		Sample		-Sample
		brevirostris		(Panthera		(n=26)
				<i>tigris,</i> n=3)	Sangiran-	Sites:
		Lutrogale		Sites:	Tiger-	Kedung
		palaeoleptonyx		Kedung	Sample	Brubus,
				Brubus and	n=14,	Kebon
				Kebon Duren		Duren, Trinil,
Trinil H.K.	0,9 ma	Pantera tigris	lower Bapang	Trinil H.K	Homotheriu	Pitu,
				Sample	m-Sample	Bumiyayu,

		Mececyon trinilensis		( <i>Panthera</i> <i>tigris,</i> n=7) Trinil and Pitu	(Homotheriu m ultimum, n=1)	Sangiran Teguan-
		Prionailurus bengalensis			Hemimacha	(Panthera
Ci Saat	1,0 ma	Panthera sp. Megacyon merriami Lutrogale palaeoleptonyx	black clays	Ci Saat- sample ( <i>Panthera</i> <i>tigris</i> , n=2) Sites: Bumiayu	irodus- Sample (Hemimacha irodus zwierzyckii, n=1) Megacvon-	tigris n=1) Jeruk- Sample (Panthera tigris n=1)
					Sample (Megacyon merrimami, n=3)	
Satir	1,2 ma	no carnivores	black clays	1		

Table 2 Fossil material for body mass reconstruction and prey focus

710 calculation. Shown are the sample name, Collection Number (Coll. No.): NM =

711 Ned.D.v/d Mijnbouw, Museum Geologi Bandung (measurements taken from

von Koenigswald, 1933); DUB = Collectie Dubois, Naturalis Biodiversity

713 Center, Leiden; SMF/PA/F = von Koenigswald Sammlung, Senckenberg

714 Research Institute, Frankfurt; Site, anatomical element (Element), portion and

side of anatomical element (Portion), reconstructed body mass [kg] (BM);

standard deviation (sd). The Pre-Ngangdong sample is a summary of the

717 Sangiran Sample, the Ci Saat , Trinil H.K., Kedung Brubus Sample.

Sample Name (from Table 1)	Coll. No.	Site	Element	Portion	BM [kg]	sd
Ngandong-Sample	NM 5497	Ngandong	Mandibula	P3 P4 M1 sin+dex	119	
	NM 49	Ngasinan	Radius	dex	134	
	NM 504	Ngandong	Mandibula	C P3fragm. P4 M1 inf dex	147	20
	NM 13776	Ngandong	Cranium	P3 P4 M1 sup dex	151	17
	NM 1184	Ngandong	Cranium	(C) (P2) P3 P4 sup dex	163	34
	NM 1933	Watualang	Humerus	sin	189	
	NM 2811	Ngandong	Cranium	complete	194	46
	NM 2671	Watualang	Mandibula	sin	212	
	NM 9554	Ngandong	Humerus	sin	235	
	2641	Ngandong	Femur	dex	298	
Kedung Brubus- Sample	DUB 1499	Kedung Brubus	Mandibula	P4 M1 inf dex	134	
	DUB 1498	Kedung Brubus	Mandibula	(C) (P3) P4(M1) inf dex	135	
	DUB 89	Kebon Duren	Femur		69	

Trinil H.KSample	DUB 8411	Trinil	Tibia	sin	99	
	DUB 1494	Trinil	Maxilla	I2 C (P2) P3 P4 (M1) sup sin	103	3
	DUB 1495	Trinil	Maxilla	(C) (P2) P3 P4 sup dex	103	5
	NM 5	Trinil	Cranium	(M1),P4,P3,(P2),(C),(I) sin M1 fragm.	133	
	DUB 1871	Trinil	Ulna	sin	117	
	DUB 1479	Trinil	Mandibula	C P3 P4 M1 inf dex	118	
	NM 61	Pitu	Mandibula	C P3 fragm. P4 M1 inf dex	114	
Ci Saat-Sample	NM 1533a	Bumiayu	Femur	sin	93	
	1209	Bumiayu	Mandibula	C1 P3 P4 M1 inf sin	96	
Teguan-Sample	DUB 90	Teguan	Ulna	dex	86	
Jeruk-Sample	DUB 6219	Jeruk	Maxilla	P3 P4 sup dex	228	10
Sangiran-Tiger-	SMF/PA/F6683	Sangiran	Praemolar	P4 sup dex	64	
Sample	DUB 11699	Sangiran	Mandibula	P3-M1 inf sin	77	14
	SMF/PA/F 6658	Sangiran	Mandibula	P3-M1 inf dex	81	8
	SMF/PA/F 6674	Sangiran	Praemolar	P4 sup sin	103	
	SMF/PA/F 6677	Sangiran	Maxilla	P4 sup sin	108	
	SMF/PA/F 6673	Sangiran	Praemolar	P4 sup dex	109	
	CD 1168 a	Sangiran	Praemolar	P3 inf sin	125	
	F6668	Sangiran	Praemolar	P4 sup sin	126	
	F6670	Sangiran	Praemolar	P4 sup dex	138	
	CD 1168 b	Sangiran	Praemolar	P4 inf sin	139	
	CD 1168 c	Sangiran	Molar	M1 inf sin	140	
	SMF/PA/F 6660	Sangiran	Maxilla	P3 P4 sup dex	143	
	SMF/PA/F 6671	Sangiran	Maxilla	P4 sup sin	153	
	SMF/PA/F 6659	Sangiran	Maxilla	P3 sup dex	182	
Homotherium- Sample	SMF/PA/F 6676	Sangiran	Mandibula	P3-M1 inf dex	154	
Hemimachairodus- Sample	SMF/PA/F 6679	Sangiran	Mandibula	P4 M1 inf dex	130	
Megacyon-Sample	F6785	Sangiran	Molar	M1 sup sin	49	
	F6792	Sangiran	Molar	M1 sup sin	51	
	F6786	Sangiran	Molar	M1 inf dex	55	

- Table 3 Regressions for reconstruction of body masses. Shown are the
- regression number (Reg. No.), the parameter (SKL = skull length), intercept
- 721 (a), slope (b), correlation coefficient (r<sup>2</sup>), standard error of estimation (SE),
- 722 percentage standard error of estimate (%SEE) and percentage standard error
- of the estimate (%PE) after Smith (1981, 1984).

Reg. No. Para	ameter a	b	r²	р	SE	%SEE	%PE
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Reg. No.	Parameter	а	b	r²	р	SE	%SEE	%PE
F-1	SKL*	-5,983	3,25	0,878	<0,0001	0,058	14	11
F-2	P2 sup	+0,632	1,389	0,314	0,02	0,442	177	76
F-3	P3 sup	-2,128	3,187	0,915	<0,0001	0,171	48	28
F-4	P4 sup	-2,775	3,239	0,948	<0,0001	0,135	36	24
F-5	P3 inf	-1,936	3,333	0,933	<0,0001	0,175	50	29
F-6	P4 inf	-2,209	3,178	0,938	<0,0001	0,158	44	26
F-7	M1 inf	-2,281	3,151	0,954	<0,0001	0,106	28	17
Reg. No.	Parameter	а	b	r²	р	SE	%SEE	%PE
C-1	P3 inf	-2,137	3,288	0,897	<0,0001	0,039	36	26
C-2	P4 inf	-2,172	3,153	0,906	<0,0001	0,039	35	23
C-3	M1 inf	-2,436	2,775	0,919	<0,0001	0,041	32	22
C-4	M2 inf	-2,035	3,346	0,831	<0,0001	0,049	51	35

Table 4 Samples for calculation of prey mass regressions. Listed is taxon,

mean body mass (BM) taken from literature (see text), pack size (PS), pack

weight (BMxPS=PW), mean prey mass (PM), number of kills used for

728 calculation of mean prey mass (n) and studies:

1= Mech and Boitani, 2003; 2 = Ruggiero, 1991; 3 = Sillero-Zbuiri and Gottelli,

730 1995; 4 = Pienaar, 1969; 5 = Power, 2002; 6 = Karanth and Sunquist, 1995; 7

= Estes and Goddart, 1967; 8 = Corbett and Newsome, 1987; 9 = Zuercher et

732 al., 2005; 10 = Johnsingh, 1983; 11= de Azevedo and Murray, 2007; 12 =

733 Polisar et al., 2003; 13 = Kruuk and Turner, 1967; 14 = Eaton, 1974; 15 =

734 Mills et al., 2004; 16 = Mills, 1990; 17 = Caro, 1994.

<sup>735</sup> \*Zuercher et al. (2005) provide no list of prey items. We therefore provide the

mean prey mass as calculated by them.

Taxon	BM [kg]	PS	PW	PM [kg]	n	Studies
Acinonyx jubatus	50	-	-	35,91	1071	4,13,14,15,16,17

Taxon	BM [kg]	PS	PW	PM [kg]	n	Studies
Panthera leo	197	-	-	238,36	2840	1,4,5,16,
Panthera onca	71	-	-	15,93	61	12,11
Panthera pardus	59	-	-	48,22	1088	4,6,10,13,16,
Panthera tigris tigris	179	-	-	401,4	157	6,10
Puma concolor	53	-	-	13,24	23	12
Canis lupus	49	9,5	465,5	251,6	248	1
Canis lupus dingo	12,5	3,3	41,25	12,7	348	8
Canis simensis	15,25	1	15,25	0,3	362	3
Cuon alpinus	15	10,5	157,5	33,4	395	6, 10
Lycaon pictus	27	11,5	310,5	44,8	859	4, 7,13
Speothos venaticus	6,5	4,5	29,25	2,2	u*	9

- Table 5 Regressions for calculation of prey mass. Shown are intercept (a),
- slope (b), correlation coefficient (r), standard error of estimation (SE),

741 percentage standard error of estimate (%SEE) and percentage standard error

of the estimate (%PE) after Smith (1981, 1984).

Nr.	Sample	Intercept	slope	r²	р	SE	%SEE	%PE
P-1	Felidae	-2,0795	1,9831	0,7922	0,017	0,138	86	60
P-2	Canidae	-2,082	1,634	0,895	0,004	0,377	138	68

- Table 6 Reconstructed body masses of the fossil carnivores. Shown are
- sample name (see Table 2), minimum reconstructed body mass (BM<sub>min</sub>),
- mean body mass (BM<sub>mean</sub>), maximum Body mass (BM<sub>max</sub>) and the range of
- statistical errors: minimum reconstructed body mass %PE (BM<sub>min%PE</sub>) and
- 748 maximum reconstructed body Mass + %PE (BM<sub>max%PE</sub>).

	BmminPE		BM mean	BM max	BM max
Sample name	% [kg]	BM min [kg]	[kg]	[kg]	%PE [kg]
Megacyon	38	49	52	55	68
Hemimachairodus	61	24	130	308	209
Homotherium	17	83	154	178	255
Tiger Samples					
(Panthera tigris)					
pre-Ngandong	49	64	114	182	233
Sangiran	49	64	121	182	233
Ci Saat	63	81	87	93	113
Trinil H.K.	68	96	110	133	171
Kedung Brubus	54	69	113	135	171
Ngandong	79	119	184	298	363
Teguan	63	86	86	86	109
Jeruk	159	228	228	228	292

- 749
- 750 Table 7 Competitors and the reconstructed prey mass spectra. The focus
- classes are framed and the percentage contribution of each focus class to the
- complete focus area is displayed.

Competitor	3a	3b	3c	4a	4b	4c	5a
Homotherium ultimum			25	25	25	25	
Machairodus zwierzyckii		25	25	25	25		
		1				1	
Megacyon merriami	20	20	20	20	20		
						1	
Sangiran liger	20	20	20	20	20		
Ci-Saat Tiger		33,33	33,33	33,33			
Trinil H.K .Tiger		25	25	25	25		
Kedung Brubus Tiger		25	25	25	25		
pre-Ngandong Tiger	20	20	20	20	20		
Ngandong Tiger		16,67	16,67	16,67	16,67	16,67	16,67

- Table 8 Results of unpaired t-test between mean body mass of the Sangiran
- sample and the Ngandong sample executed by Graph Pad "Quick Calcs"
- 756 http://www.graphpad.com/quickcalcs

P value and statistical significance:						
The two-tailed P value equals 0.0073	By conventional criteria, this difference is considered to be statistically significant.					
Confidence interval:						
The mean of Group One minus						
Group Two equals	-55.14					
95% confidence interval of this	From -94.59 to					
difference	-15.69					
Intermediate values used in calcula	ations:					
standard error of difference	19.520					
t	2.8246					
df	40					
Group	Sangiran	Ngandong cample				
Gloup	sample	riganuong sample				
Mean	129.06	184.20				
SD	53.92	53.74				

SEM	9.53	16.99
Ν	32	10

757 Figures

758 Figure 1

Large carnivore assemblages from several faunal levels of the Pleistocene of

Java. Shown are only carnivores weighing more than 10 kg. The Sangiran v.K

assemblage reflects taxa collected in the Sangiran dome by von Koenigswald

and excludes finds by the team of Sémah (Bouteaux et al., 2007).

Figure 2

764 Map of Java and the sites of the fossil assemblages.

765

Figure 3

767 Reconstructed body masses of the Pleistocene tiger samples, the

768 Machairodonts and *Megacyon merriami* in Java. Each reconstructed body

mass value is displayed as a black dot for the samples. The sample name is

provided at the x-axis. Composition of the samples is given in Tables 1 and 2.

771

Figure 4

Niche overlap by hypercarnivorous carnivores from Sangiran with tigers.

Shown are the different tigers and their niche overlap with *Homotherium* 

175 ultimum, Hemimachairodus zwierzyckii and Megacyon merriami.

\* The samples Sangiran and pre-Ngandong have the same PMS and thus the

same niche overlap in percent.

<sup>778</sup> \*\* the samples Ci Saat, Trinil H.K. and Kedung Brubus have the same PMS

and thus the same niche overlap.

780

781 Figure 5

- 782 Niche overlap by tigers with other hypercarnivores from Sangiran. Shown are
- 783 Homotherium ultimum, Hemimachairodus zwierzyckii and Megacyon merriami
- and their niche overlap with tigers in percent. \* The samples Sangiran and
- 785 pre-Ngandong have the same PMS and thus the same niche overlap.
- <sup>786</sup> \*\* The samples Ci Saat, Trinil H.K. and Kedung Brubus have the same PMS
- and thus the same niche overlap.

Cover Letter, for Editor only

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Manila, 19 July 2015

Elsevier Palaeogeography, Palaeoclimatology, Palaeoecology Amsterdam Netherlands

### Dear editors,

We hereby submit our manuscript with the title "Niche overlap and competition potential among tigers (*Panthera tigris*), sabertoothed cats (*Homotherium ultimum*, *Hemimachairodus zwierzyckii*) and Merriam's Dog (*Megacyon merriami*) in the Pleistocene of Java".

We investigated the role of competition in a case of dramatic body mass increase in Late Pleistocene tigers. During the Late Pleistocene, tigers with a body mass in excess of 300 kg inhabited Java, whereas tigers from the Early and Middle Pleistocene of Java range in body mass between 63 until 121 kg. The question is what drove this size increase in tigers?

In order to answer this question, we estimated body mass of tigers from all periods, and of its potential competitors. Based on body mass and new regressions for both felids and canids derived from extant species, we calculated prey mass spectrum and niche overlap. Niche overlap was then used as proxy for degree of competition. We found that Merriam's dog, a very large wolf-like canid, had a 100% niche overlap with tigers. We concluded that in order to minimize competition, tigers increased in size whereas Merriam's dog decreased in size (ecological character displacement). The sabertoothed felids, which also had a considerable overlap, went extinct.

On behalf of all authors, Sincerely yours,

Rebekka Volmer

Figure1





50 km





Figure4



Figure5



Highlights

- Late Pleistocene tigers of Java belong to the largest known tigers
- Shifts in body masses of tigers are probably caused by competition.
- Tigers on Java had highest competition potential with Merriam's Dog.
- *Homotherium ultimum* had the lowest competition potential with tigers.
- New regressions for body mass and prey mass reconstruction for large carnivores were calculated.

1 Supplementary data

2 Material

3 A variety of studies provides regression terms for parameters like upper M1 length 4 and long bones for the estimation of body mass (van Valkenburgh, 1990; Anyonge, 5 1993; Hemmer, 2001; Christiansen and Harris, 2005). Skull length and lower M1 6 length are used as well. Teeth other than M1 have not yet been studied for correlation with body mass. We were therefore unable to reconstruct a large portion 7 8 of the specimens in the collections (Hertler and Volmer, 2008). Many fossil samples 9 only encompass dental material, for example, samples from the Sangiran dome. 10 The Ngandong sample includes a tiger cranium based on which Hertler and Volmer 11 (2008) reconstructed a large body mass of 470 kg using the felid specific regression 12 of van Valkenburgh (1990). Because body size and skull length of tigers varies 13 hugely (Mazák, 1981), it is advisable to calculate a species-specific regression term based on tigers only (Panthera tigris) of Mazák's dataset (Mazák, 1981). In our 14 15 present study additional regressions were calculated to include additional tooth positions in the reconstruction of body masses for Felidae and Canidae . 16 17 Furthermore, a tiger specific regression term is calculated. 18 19 Body mass regressions

A sample of extant skeletons, representing 31 felid and 16 canid species, of the Senckenberg Collection were measured for the regression terms. Only fully erupted, complete, permanent teeth are measured. In order not to confuse intra- and interspecific allometry, averages of each species with multiple individuals were used for statistical analysis. For some taxa and parameters larger samples than available in the Senckenberg Institute were already measured and published by Schmid (1940). In these cases the samples are used instead. 27 The length of the teeth is defined as the largest mesio-distal length and was

28 measured with digital vernier calipers. For each taxon and parameter the means of

29 the measurements is calculated. The mean tooth length and the mean body mass,

30 taken from literature are defined as a pair of values.

31 For the regression for tiger skull length the data from Mazák (1981) were used. Here

32 the pair of values is the minimum and maximum skull lengths and body masses,

33 including the sex. For example: the maximum body mass for male Sumatra Tiger is

34 paired with the maximum body mass for male Sumatra Tiger. This method gives a

- 35 higher sample size.
- 36 The samples are shown in Supplementary Tables 1–3. The linear regressions for

37 reconstruction of body masses are computed by Systat 12 using tooth/skull length as

38 known variable and body mass as the unknown variable.

- 39
- 40 Regressions for Reconstruction of Body Masses
- 41 The calculated regressions for body mass reconstruction are shown in Table 5. A
- 42 statistically relevant correlation with body mass is present for all parameters (P</=
- 43 0,02). All parameters show a strong correlation ( $r^2 > 0,878$ ), except for the upper P2
- 44 length. The percentage estimation errors vary between 14–177%, the %PE between
- 45 11–76 %.
- 46

## 47 Supplementary Tables

48

Supplementary Table 1 Sample for calculation the regression for body mass reconstruction of
skull length of *Panthera tigris* (Regression F-1). Listed are the subspecies, sex, minimal body mass
(BM min), maximal body mass (BM max) and minimal and maximal skull length (SKL min, SKL max).
BM min and SKL min of one sex of a subspecies is used as a pair of values and BM max and SKL
max respectively. All data taken from Mazák (1981).

Subspecies	Sex	BM min	BM max	SKL min	SKL max
P.t. altaica	М	180	306	341	383
P.t. altaica	F	100	167	279	318
P.t. amoyensis	Μ	130	175	318	343
P.t. amoyensis	F	100	115	273	301

Subspecies	Sex	BM min	BM max	SKL min	SKL max
P.t. balica	M	90	100	295	298
P.t. balica	F	65	80	263	269
P.t. corbetti	Μ	150	195	319	365
P.t. corbetti	F	100	130	279	302
P.t. sondaica	М	100	141	306	349
P.t. sondaica	F	75	115	270	292
P.t. sumatrae	М	100	140	295	335
P.t. sumatrae	F	75	110	263	294
P.t. tigris	Μ	180	258	329	378
P.t. tigris	F	100	160	275	311
P.t. virgata	Μ	170	240	316	369
P.t. virgata	F	85	135	268	305

55 Supplementary Table 2 Samples for calculation of body mass regressions based on dental

56 parameters for Felidae. Listed are taxon, mean body mass (BM) [kg], the tooth length [mm] (P2 sup,

57 P3 sup, P4 sup, P3 inf, M1 inf) and abbreviation of the regression (F-2 – F-7), n= sample size. Source

58 of data (S) :1= parameters taken from specimens housed at Research Institute Senckenberg, 2 =

59 Schmid (1940), 3 = Hooijer (1947), 4 = Mazlaghani (2005), 5 = Hemmer (1971), 6 = Brongersma

60 (1935).

#### 61 \* = only female individuals, mean body mass of females is used

		F-2		F-3		F-4		F-5		F-6		F-7		
	BM	P2		P3		P4		P3		P4		M1		
Taxon	[kg]	sup	n	sup	n	sup	n	inf	n	inf	n	inf	n	S
Leopardus														
pardalis	9,9	4,53	1	11,13	1	12,27	1	9,55	1	12,08	1	12,71	1	1
Neofelis														
nebulosa	15,5	2,45	1	12,6	16	18,5	16	8,5	16	13,8	16	14	16	1,2
Panthera														
leo	197	8,49	2			33,59	13	15,76	10	23,96	12	24,65	12	1
Panthera														
onca	71	6,94	9			27,18	12	14,23	12	19,76	12	20,56	13	1
Panthera														
tigris														
balica*	72,5	18,98	1	19,1	1	30,05	1					21,57	1	1
Panthera														
tigris						00.4								
corbetti	147,5	6,9	1	20,9	1	33,4	1	21,8	1			26	1	4
Panthera	470	F 70		00.0		05.05		10.5		00 F		00.0		~
tigris tigris	179	5,72	1	22,3	u	35,65	u	16,5		23,5		26,9	u	3
Prionailurus	0.05	4.04		0.04				0.0		0.05				1.0
planiceps	2,25	4,34	1	8,34	1	11,4	1	6,2		8,35	u			1,6
Prionaliurus	0	2.00	4	0.64	4	10.00	1	6.04	4	0.20	4	10 57	1	4
Viverninus	9	2,69	1	0,04	1	13,00	1	0,01	1	9,30	1	10,57	1	1
tioris														
altaious	202	7 92	1	22.01	2	22.15	4	17 29	2	22.04	4	25.02	4	12
Acinony	203	7,05	1	23,01	3	55,15	4	17,50	3	23,94	4	23,92	4	1,5
iubatus	50	2.8	з	13.04	6	21 51	6	13.86	4	15 1	6	18.2	6	з
Panthera	50	2,0	5	10,04	0	21,01	0	10,00	-	10,1	0	10,2	0	5
nardus	59	5	82	16.3	113			11.8	94	173	aa	18	104	2
Panthera	55	5	02	10,0	110			11,0	54	17,5	55	10	104	2
pardus														
fusca	45.5	5.2	9	16.4	12	25.1	12	11.7	12	17.8	12	17.8	11	2
Panthera	.0,0	0,2	Ű	,.		_0,:		,.		,e		,e		-
pardus														
melas	35	3.7	12	14.8	23	22.2	22	10.5	22	15.3	24	16	25	2
Panthera		-,-		,e		,_		,.		,.				
pardus														
orientalis	36,5	5,4	1							18,75	1	18,65	1	2
Puma				1										
concolor	53	5,5	6	15,23	13	21,53	15	11,79	11	14,76	13	17,81	12	1
Uncia uncia	45	6,3	28	15	30	24,1	29	12,3	29	16,6	29	18	28	2
Panthera	107,5	8,3	13	20,8	13	31,35	13	14,55	13	21,95	13	23,74	13	1

tigris													
l vnx lvnx	19.85	x	11.9	20	18.8	22	99	29	12.2	29	15.7	31	2
Catopuma	10,00			20	10,0	~~~	0,0	20	12,2	20	10,7		-
temincki	11 75						81	3					1
Felis chaus	10	ł					7.75	4	10	4	10.55	4	5
Panthera		1					.,						
leo*	152				32,87	5	16,28	3					1
Panthera													
pardus													
(Africa)	59				24,5	109							2
Panthera													
tigris													
amoyensis	137,5		21,9	5	33,5	5	16,15	5	22,65	5	24,75	5	3
Panthera													
tigris			~~~~										
sondaica	108		20,65	8	32,31	12	14,9	8	21,74	14	24,18	15	1,3
Pardotelis													
badia	4			<u> </u>	<u> </u>		5,9	<u> </u>	7,8	u	ļ	<u> </u>	6
Pardotelis	25		7 40		44.04								
marmorata	3,5	<b>├</b> ───	7,48	1	11,84	1	5,55		8	u	<u> </u>	<u> </u>	6
Pardotelis	44 75		0.05	2	10.40	2			10.00	_	44.04	2	4
temminckii	11,75	<b>├</b> ───	9,95	3	16,42	3	───		10,66	3	11,84	3	1
Prionaliurus	E 1E		5 00	1	0.07	12	5		6 5	l	7.66	12	1.6
Derigaierisis	5,15	<u>├───</u>	5,99		9,97	13	5		6,5	u	1,00	13	1,0
Phonaliurus	1.5						4		5 25				6
Profolio	1,5	<u> </u>					4	-	5,35	u	<u> </u>	<u> </u>	0
PIUlelis	11 75		9.18	3	15 49	3	7 25	3	10.00	3	11 27	3	1
aulala	11,75		3,10	5	13,43	5	1,25	5	10,03	5	11,21	5	

63 Supplementary Table 3 Samples for calculation of body mass regressions based on dental

64 parameters for Canidae. Listed are taxon, mean body mass (BM) [kg], tooth length [mm] (P3 inf, P4

65 inf, M1 inf), and abbreviation of regression (C-1 – C-3), n= sample size.

Taxon	BM	C-1		C-2		C-3	
		P3 inf	n	P4 inf	n	M1 inf	n
	[kg]	[mm]		[mm]		[mm]	
Alopex lagopus	3,685	7,35	5	8,1	5	12,72	5
Canis adustus	9,65	7,4	4	8,14	4	13,74	4
Canis aureus	8,15	8,95	5	10,05	5	18	5
Canis latrans	11,75	11,12	2	12,16	2	20,74	2
Canis lupus arctos	49	13,73	11	15,47	13	28,26	13
Canis lupus dingo	12,5	10,6	8	12,38	8	21,12	9
Canis mesomelas	9,25	8,51	8	9,95	7	16,56	5
Canis simensis	15,25	9,42	1	10,38	1	18,56	1
Cerdocyon thous	5,7	7,75	4	8,61	4	14,92	4
Chrysocyon brachyurus	25	11,58	5	12,92	5	21,52	5
Cuon alpinus	15	9,8	2	12	3	21,63	3
Lycaon pictus	27	11,57	6	13,19	7	24,52	7
Speothos venaticus	7,715	8,54	1	8,99	1	14,36	1
Urocyon cinereoargenteus	3,75	5,9	3	7,3	3	12,31	3
Vulpes bengalensis	2,5	6,34	1	7,13	1	10,71	1
Vulpes zerda	1,25	5,11	3	5,21	3	8,46	3

66

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