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1 *American Journal of Primatology*

2 **Gross intestinal morphometry and allometry in primates**

3 **Running title: Primate intestinal length**

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24 **Abstract**

25 Although it is generally assumed that among mammals and within mammal groups, those
26 species that rely on diets consisting of greater amounts of plant fiber have larger
27 gastrointestinal tracts (GIT), statistical evidence for this simple claim is largely lacking. We
28 compiled a dataset on the length of the small intestine, caecum and colon in 42 strepsirrhine,
29 platyrrhine and catarrhine primate species, using specimens with known body mass (BM).
30 We tested the scaling of intestine length with body mass, and whether dietary proxies
31 (percentage of leaves and the diet quality index) were significant covariates in these scaling
32 relationships, using two sets of models: one that did not account for the phylogenetic
33 structure of the data, and one that did. Intestine length mainly scaled geometrically at
34 exponents that included 0.33 in the confidence interval; Strepsirrhini exhibited particularly
35 long caeca, while those of Catarrhini were comparatively short. Diet proxies were only
36 significant for the colon and the total large intestine (but not for the small intestine or the
37 caecum), and only in conventional statistics (but not when accounting for phylogeny),
38 indicating the pattern occurred across but not within clades. Compared to terrestrial
39 Carnivora, primates have similar small intestine lengths, but longer large intestines. The data
40 on intestine lengths presented here corroborate recent results on GIT complexity, suggesting
41 that diet, as currently described, does not exhaustively explain GIT anatomy within primate
42 clades.

43

44 **Keywords** primate, anatomy, digestive tract, diet, phylogeny

45 **Introduction**

46 Of the various aspects of mammalian biology where a link between form and function has
47 been investigated, the connection between diet and the morphology of the digestive tract has
48 received considerable attention (Chivers and Langer, 1994). Illustrations of the digestive
49 tracts of herbivores and carnivores make it evident that the digestive tracts of herbivores are
50 more complex (Stevens and Hume, 1998) due to the less digestible nature of their natural
51 diets. Though statistical evidence is lacking (Lavin, Karasov, Ives, Middleton and Garland,
52 2008; Smith, Parker, Kotzé and Laurin, 2017), it is usually thought that the intestines of
53 herbivorous mammals are longer than those of carnivores (Orr, 1976). When considering
54 gastrointestinal complexity across all eutherian mammal clades, however, there is a less
55 clear-cut relationship between intestine morphology and diet. In particular, while complex
56 hindguts are typically associated with (but not obligatory for) herbivores, complex stomachs
57 are not necessarily linked with high-fiber herbivory, as evidenced by the morphology
58 observed in whales and myomorph rodents (Langer and Clauss, 2018).

59 Among primates, the morphology of the gastrointestinal tract, both macro- and
60 microscopically, has been suggested to be tightly linked to dietary ecology (Chivers and
61 Hladik, 1980; Martin, Chivers, MacLarnon and Hladik, 1985; Lambert, 1998). Understanding
62 aspects of the anatomy of primates' digestive systems is thought to provide information about
63 the diet to which a particular taxon may be adapted (Ferrari and Lopes, 1995) or its dietary
64 constraints, such as why a species may target or avoid certain food items (Lambert, 1998).
65 Despite the value of broad anatomical comparisons across taxa, relatively few studies have
66 taken an interspecific, phylogenetic approach to the gross anatomy of primate gastrointestinal
67 tracts. Quantitative analyses of gut morphology frequently focus on a small range of taxa
68 (e.g., Ferrari and Martins, 1992; Ferrari, Lopes and Krause, 1993 in callitrichines), and some
69 studies (e.g., Hill, 1958) provide illustrations of primate gastrointestinal tracts, but do not

70 provide measurements. The paucity of recent gross intestinal morphology research may be in
71 part related to the fact that the proportions of different components of the gastrointestinal
72 tract can vary intraspecifically with age (Langer, 1988) or with dietary shifts (Gross, Wang
73 and Wunder, 1985). Nevertheless, the use of relative dimensions can provide information
74 about the relative role of different portions of the tract in a particular subset of species
75 (Chivers and Hladik, 1980; Caton, 1998) and can inform hypotheses about adaptations to
76 different dietary regimes. The enlarged caecum and colon of some strepsirrhines, New World
77 monkeys, cercopithecines, and hominoids (Hladik, Charles-Dominique, Valdebouze, Delort-
78 Laval and Flanzky, 1971; Chivers, 1994), for example, act as fermentation chambers to aid in
79 the digestion of plant foods.

80 On the other hand, it has also been recognized that gastrointestinal form alone is not
81 sufficient to predict diet (Milton, 1987), even though general relationships between gross
82 digestive morphology (e.g., relative gut proportions) and broad dietary categories are
83 expected. Chivers and Hladik (1980) demonstrated an increase in the ratio of combined
84 stomach, caecum, and colon volume to small intestine volume across a continuum of
85 faunivores to frugivores to folivores, although there was substantial overlap between the
86 groups. When Chivers and Hladik considered the surface area of the small intestine, however,
87 no differences between trophic groups were evident. Notably, these analyses were performed
88 prior to the arrival of statistical methods that account for phylogenetic non-independence.
89 When controlling for phylogeny, Langer and Clauss (2018) did not find diet-associated
90 differences in gut complexity among primates.

91 Despite these various lines of research, quantitative data on the digestive tracts of
92 primates are scarce. In order to provide data on intestine length for a larger number of
93 primates, we used the opportunity to take photographs of the digestive tracts of a number of
94 specimens during dissections performed for a study on mammalian body composition

95 (Navarrete, van Schaik and Isler, 2011), and provide photographic evidence for the gross
96 digestive tract morphology of 32 species. Because with any absolute measure, body size will
97 be the main driver of the magnitude, we first investigate the allometric relationship of
98 intestine length with body mass. Based on the general geometric relationship between a
99 length and a volume measure, one would expect intestinal lengths to scale approximately
100 with body mass to the power of 0.33. However, three previous data compilations – on
101 mammals in general (Lavin et al., 2008), on mammalian carnivorans (McGrosky, Navarrete,
102 Isler, Langer and Clauss, 2016) and on ruminants (McGrosky et al., 2019) – unexpectedly
103 yielded higher exponents. The explanation for this phenomenon was, to our knowledge, first
104 proposed by Woodall & Skinner (1993), who suggested that animals should evolve so that
105 their intestinal surface retains a geometric or metabolic scaling (i.e., at an exponent between
106 0.67 and 0.75), but that the diameter of the intestine should scale less-than-geometrically to
107 maintain short diffusion distances, and that hence, to compensate, intestinal length should
108 scale more-than-geometrically. Therefore, a corresponding scaling was expected in primates
109 as well.

110 As a second step, we tested whether proxies of the natural diet influenced the scaling
111 relationships with body mass. Following broad mammalian trends and Chivers and Hladik
112 (1980), we expect both small and large intestine length to increase along a dietary continuum
113 from insectivores to frugivores to folivores.

114

115 **Methods**

116 During the data collection for a different study (Navarrete et al., 2011), a large number of
117 mammalian specimens originating from various sources were dissected by AN; some
118 additional specimens were contributed by MC. Most specimens derived from zoological
119 collections. The specimens were either dissected directly after death or, in the majority of

120 cases, after storage by deep-freezing and thawing. Body mass (BM) was recorded, and the
121 gastrointestinal tracts were dissected by removing mesenteries. The intestinal tract was laid
122 out according to common practice for anatomical documentation (Stevens and Hume, 1995),
123 placing the intestine as a straight line without stretching, and photographs (with scale) were
124 taken for later measurement. For measurements, the caecum served as the defining border
125 between the small and the large intestine. Length of the different sections was determined
126 using the polyline VOI tool of the MIPAV biomedical imaging software (version 14.0,
127 National Institutes of Health), which facilitated tracing the length of the intestine directly on
128 the digital photo and converting pixel length into centimeters.

129 For some large-sized specimens, the intestine was cut into several pieces prior to
130 photography. Photographs of these specimens were not manipulated to create the impression
131 of an uninterrupted intestine. The only photo manipulations included the deletion of
132 background and, in some cases, adjustment of the brightness and color of the images and a
133 mirroring to achieve a consistent position of the stomach.

134 Additional data were collected from the literature, but only if the original BMs of the
135 specimens investigated were reported together with the length measurements; this led to the
136 exclusion of many published intestine length measurements. When several specimens were
137 available for a species, averages for both BM and the respective measurements were
138 calculated; in doing so, we always ensured the averaged BM matched the averaged gut
139 section measure.

140 Data on the natural diet of the investigated species, as the percentage of leaves in the
141 overall diet (as generally the diet items most requiring microbial fermentation) and the diet
142 quality index (to account not just for one but the range of diet items), were taken from a
143 literature collection based on three publications (Van Woerden, van Schaik and Isler, 2010,
144 2014; Van Woerden, Willems, van Schaik and Isler, 2012), supplemented with additional

145 information (for a full set of sources, see Table S1). The diet quality index represents a sum
146 of each dietary item, multiplied by a factor that is higher for more digestible diet items (8 for
147 insects, 5 for fruits/seeds/flowers, 3 for young leaves, 1 for mature leaves) and is weighted
148 for the proportion of these items in the diet (Van Woerden et al. , 2010). All species-specific
149 data are given in Table 1.

150 Allometric relationships were investigated using linear regressions on log-transformed
151 data according to $\text{Length} = a \text{BM}^b$, with 95% confidence intervals for parameter estimates,
152 for the whole dataset and for individual clades. Additionally, the same analysis was
153 performed with clades as cofactors, and the clade x BM interactions, to directly test for
154 differences between the clades. Proxies of the natural diet were added to the linear
155 regressions on log-transformed data as covariates (only one proxy per model, i.e. either the
156 percentage of leaves, or the diet quality index). Initially, models including the BM x diet
157 proxy interaction were also included, but because there were no significant interactions,
158 results are only presented here for models without interactions. Analyses were performed
159 using Generalized Least Squares (GLS) and Phylogenetic Generalized Least Squares (PGLS),
160 using a consensus phylogenetic tree downloaded from the 10kTrees Project (version 3;
161 Arnold, Matthews and Nunn, 2010). The phylogenetic signal (Pagel's λ) was estimated using
162 maximum likelihood (Revell, 2010). λ can vary between 0, indicating no phylogenetic signal,
163 and 1, indicating a strong phylogenetic signal and that similarity among species scales in
164 proportion to their shared evolutionary time (Pagel, 1999; Freckleton, Harvey and Pagel,
165 2002). Statistical tests were performed using the package caper (Orme et al., 2010) in R
166 2.15.0 (Team, 2011). The significance level was set to 0.05.

167 Finally, the data of primates were compared visually to data from terrestrial Carnivora
168 including canids, mustelids, ursids, mephitids, procyonids, felids, hyenids, herpestids and
169 viverrids, but excluding pinnipeds (McGrosky et al., 2016).

170

171 **Results**

172 Among the Strepsirrhini, lemurs appear to have particularly long caeca (Fig. 1) compared to
173 the gummivorous cheirogalids (Fig. 2). The latter have smaller caeca, although their caeca
174 still appear proportionally larger than those of the gummivorous callitrichids (Fig. 3). The
175 non-gummivorous cebids studied here all have similar digestive tracts (Fig. 4). Atelids have
176 an unremarkable caecum that appears part of a large proximal colon (Fig. 5), whereas the
177 caecum of the pitheciids is more distinct (Fig. 5). Among the Catarrhini, colobines have a
178 voluminous forestomach, as well as a distinct caecum and an enlarged proximal colon (Fig.
179 6). The hindgut of many non-colobine Catarrhini is voluminous and haustrated, with a short,
180 broad caecum (Fig. 7-10). In the apes, the caecum bears a clearly visible, vermiform
181 appendix (Fig. 10).

182 When using the complete gastrointestinal morphology dataset, as well as Strepsirrhini-
183 and Platyrrhini-specific datasets, there was a strong, significant phylogenetic signal,
184 indicating that the data scatter is guided by the phylogenetic structure in the datasets.
185 However, this strong phylogenetic signal was generally not observed within Catarrhini (Table
186 2). Across all species, the scaling exponent typically included 0.33 in the 95% confidence
187 interval, indicating geometric scaling (Table 2). The 95% confidence intervals for the
188 allometric factors (a) and scaling exponents (b) generally overlapped between the clades,
189 with the exception of both the factor and exponent of the caecum, which were higher in the
190 Strepsirrhini (Table 2). This observation was corroborated by the models that tested the
191 scaling including the clade and clade x BM interactions, where the Strepsirrhini x BM
192 interaction was significant both in GLS ($P = 0.009$) and PGLS ($\lambda = 0.73$; $P = 0.017$),
193 indicating a steeper scaling of the caecum with BM for this clade (Fig. 11). Additionally,
194 although the corresponding 95% confidence intervals for a overlapped, the analyses for small

195 intestine length and total intestine length indicated significantly smaller values for Platyrrhini
196 (small intestine: GLS $P = 0.015$; PGLS $\lambda = 0.73$, $P = 0.018$; total intestine: GLS $P = 0.032$;
197 PGLS $\lambda = 0.80$, $P = 0.035$; in all cases, with non-significant interaction terms; Fig. 11).

198 While the scaling exponent of GLS models was generally very similar to that of PGLS
199 models (Table 2 and S2), the GLS scaling exponent for the caecum in the complete dataset
200 was very low (due to the long caeca of small-bodied strepsirrhines) and excluded geometric
201 scaling in the confidence interval (0.16; 95%CI 0.04-0.29; Fig. 11). However, PGLS analyses
202 indicated a generally geometric scaling of the caecum (0.41; 95%CI 0.25-0.56; Table 2).

203 Neither the percentage of leaves in the diet nor diet quality index exhibited a significant
204 effect on intestinal length in any of the models, including models of caecum length (P always
205 > 0.05). The only exceptions to this trend included significant GLS models of colon and total
206 large intestine length for the percentage of leaves dataset and the diet quality index (GLS:
207 colon – leaves $P = 0.045$; colon – diet index $P = 0.008$; large intestine – leaves $P = 0.066$;
208 large intestine – diet index $P = 0.011$). In PGLS, these dietary effects were no longer
209 significant, indicating that variation in large intestine length with diet occurred mainly across
210 and not within clades (PGLS: colon – leaves $P = 0.402$; colon – diet index $P = 0.075$; large
211 intestine – leaves $P = 0.648$; large intestine – diet index $P = 0.231$). Using the diet quality
212 index and the relative colon length (in $\text{cm}/\text{BM}^{0.33}$), Figure 12 illustrates how the phylogenetic
213 structure of the dataset explains the colon – diet index relationship: within clades, there is
214 little correlation between the measurements, and any trend within Strespirrhini or Platyrrhini
215 is due to a particular genus, not due to similar variation in several genera; the position of the
216 clades, however, suggests the expected negative relationship. For a more detailed graph
217 including the phylogenetic tree and the distribution of these two measures across it, see Fig.
218 S1.

219 When compared to terrestrial Carnivora, the total intestines of primates are generally
220 slightly longer (Fig. 11). This holds particularly true for the colon, and the caecum of the
221 Strepsirrhini. By contrast, the caecum of the Catarrhini was of a magnitude also observed in
222 many terrestrial Carnivora.

223

224 **Discussion**

225 Using what is to our knowledge the most comprehensive dataset on primate intestine lengths
226 directly linked to the body mass of the same specimens, the present study largely
227 corroborates well-known features of different primate groups and in part confirms broad
228 macroanatomical concepts about differences in intestinal length between carnivores and
229 herbivores. However, it also demonstrates that, in primates, correlations of length measures
230 of the different intestinal sections with the natural diet **may not be** as clear-cut as is often
231 assumed.

232 **The photographic evidence provided can be compared to drawings and reports from the**
233 **literature. While we do not want to dwell on this in detail here, some points of interest shall**
234 **be mentioned. Fisher (2000) explained a reliable classification of species with respect to**
235 **whether they have a true ‘appendix’ or not requires more than visual inspection, including**
236 **histology, and ontogenetic series. Several species that have been classified as having an**
237 **appendix in comparative datasets (e.g., Smith et al., 2017), including *Eulemur* spp. (Fig. 1)**
238 **and *Callithrix* spp. (Fig. 3), did not have a vermiform structure that was clearly**
239 **distinguishable. For *Papio hamadryas*, a variable disposition with respect to the presence of**
240 **an appendix has been assumed (Smith et al., 2017), with our specimen not showing**
241 **macroscopic evidence of a vermiform structure (Fig. 9). With respect to *Varecia* spp. or**
242 ***Lemur catta*, it is evident that the macroscopic appearance could be interpreted both ways**

243 (Fig. 1), and that further investigations as recommended by Fisher (2000) would be required
244 for a reliable assessment.

245 A series of limitations applies to the present study. Our own and the literature data
246 represent a mixture of measurements taken from freshly dissected specimens and specimens
247 dissected after frozen storage and thawing. While we do not think that this has a relevant
248 effect on macroscopic length measurements, readers that question this assumption should
249 therefore consider the results with extra caution. In our view, it needs to be mentioned that
250 length measurements of intestinal structures should rather be considered as indicating
251 magnitudes, and not precise data (hence, we give no decimals in Table 1). When laying out
252 intestines for length measurements, details such as the dryness of the organ and the
253 smoothness of the surface used for measuring can influence whether an unintentionally
254 stretched organ can contract again or retains the stretched disposition, for example. A further
255 kind of caveat that needs to be mentioned refers to the use of a consensus tree (here, from the
256 10kTrees Project) rather than testing a whole distribution of trees. The standpoint, adopted
257 intuitively in the present study, that absence of a signal when using the consensus tree is
258 meaningful, could of course be questioned, and the data could be explored using the whole
259 distribution of trees available.

260 However, the main limitations of the present study are its reliance on mainly captive
261 specimens, and the low sample size for the majority of the included species that also
262 precludes tests of ontogenetic changes. Freckleton, Harvey, & Pagel (2002) demonstrated
263 that at lower species numbers below 20, a phylogenetic signal may not be always detected
264 reliably, whereas a wrong detection of a phylogenetic signal is rare. In our analyses, this was
265 also reflected in the finding that confidence intervals for λ could often not be calculated
266 (Table 2). Among the Strepsirrhini (n=11) and Platyrrhini (n=14), the limited number of
267 species might have been responsible for the finding that small intestine length and colon

268 length (and hence also large intestine and total intestine length) showed a phylogenetic
269 signal. This phylogenetic signal indicates that variation in these measures and clades was
270 mainly an effect across, and not within, the included taxonomic subgroups. Notably, this did
271 not apply to the Catarrhini, for which we obtained a larger sample size (n=17). In contrast,
272 the fact that no phylogenetic signal was evident in strepsirrhine caecum length suggests that a
273 long caecum that increases in length with body mass is a common feature across various
274 strepsirrhine subgroups. For more confidence in the phylogenetic signal of intestinal length
275 measurements, a larger species sample would be required. This would also help to clarify if
276 primates are really different from carnivores, ruminants or mammals in general (Lavin et al.,
277 2008, McGrosky et al., 2016, 2019), which all show a more-than-geometrically scaling of
278 intestine length, possibly to achieve geometrical constancy of absorptive intestinal surface
279 while keeping diffusion distances in the intestine small by increasing the intestinal diameter
280 at less-than-geometrically scaling (Woodall and Skinner, 1993). The large 95% confidence
281 intervals for the scaling exponents in the present study do not allow a clear answer to this
282 question.

283 Another limitation becomes evident given the absence of a clear dietary signal in
284 intestinal length measurements, and is inherent to our study design: intestinal length is
285 possibly poorly correlated with diet. Other measures, notably the volume or the actual gut fill
286 (Clauss, Schwarm, Ortmann, Streich and Hummel, 2007; Müller et al., 2013), are most likely
287 more meaningful correlates with diet, as mammalian herbivores generally have more
288 voluminous (yet not necessarily longer) intestines. Chivers and Hladik (1980) addressed this
289 fact by using intestinal surface area rather than length for some of their analyses. Notably,
290 they did not find a significant difference in small intestine surface area between diet groups,
291 and they did not report results for the large intestine or caecum in isolation, as done in the
292 present study, where any dietary signal that derives from the anatomy of the stomach and

293 forestomach was excluded. The often-cited patterns between gut morphology and diet
294 described by Chivers and Hladik (1980) always represent a composite signal in which the
295 volume or surface area of the (fore)stomach and the large intestine are summed up before
296 evaluation. However, even using this composite signal, their study revealed substantial
297 overlap between faunivores, frugivores and folivores. Additionally, affiliation to dietary
298 groups varied, depending on whether anatomical surface or volume measurements were used;
299 while colobine monkeys were exclusively grouped as folivores in terms of volume measures
300 (Fig. 20 in Chivers and Hladik, 1980), they were also depicted within the range of frugivores
301 in terms of surface measures (Fig. 18 of their study).

302 Nevertheless, the present study provides evidence for the functional relationship between
303 colon or large intestine length and diet, as postulated previously (Hladik, 1978; Chivers,
304 1994). Similar to the findings on gastrointestinal complexity of Langer and Clauss (2018),
305 GLS models revealed a significant relationship between diet and large intestine length.
306 However, in both Langer and Clauss (2018) and the present study, these effects were not
307 significant when accounting for the phylogenetic structure of the data using PGLS. This
308 indicates that these effects do not occur within the different primate clades and therefore
309 cannot be considered convergent between them in the respective datasets. Rather, the pattern
310 occurs at deeper nodes of the phylogeny in each dataset, to the extent that clades as a whole
311 differ in both, the GIT and the diet measure. The lack of significance in PGLS, or, in other
312 words, the lack of evidence for dietary convergence, should not encourage the interpretation
313 of a lack of a functional relationship. The relationship is just not evident *within* clades, but
314 only *across* the clades included in this study. This could lead to the traditional interpretation
315 often applied to non-significant findings when accounting for phylogeny, i.e. clade-specific
316 characteristics other than diet (represented by ‘phylogeny’) led to the evolution of longer or
317 more complex large intestines in those taxa that also ingest more leaves. Alternatively, it

318 suggests that clade diversification in primates followed, to a large extent, dietary niche and
319 concomitant intestinal adaptations, but that diet diversification within clades left less reliable
320 marks on intestinal anatomy. Reporting only one result, such as – in the case of a significant
321 phylogenetic signal – the PGLS result while ignoring the GLS result, as sometimes
322 recommended in the biological literature (Freckleton, 2009), would leave this observation
323 unrecognized.

324 With respect to length, primates do not appear to differ from terrestrial carnivores for the
325 small intestine where auto-enzymatic digestion occurs. However, as also described by Martin
326 (1990) and Lambert (1998), primates generally have a longer large intestine (a site of allo-
327 enzymatic digestion), which coincides with their generally higher reliance on diets that
328 contain plant cell walls compared to carnivores. The difference in the intestinal segment by
329 which primates achieve a longer large intestine resembles the pattern observed between other
330 large (e.g., perissodactyl) and small (e.g., rodents and lagomorphs) hindgut-fermenting
331 herbivores (Stevens and Hume, 1998); while smaller species typically have particularly large
332 caeca and are often called ‘caecum fermenters’, larger species also possess a voluminous
333 colon for allo-enzymatic digestion. This pattern is reflected in the differences between the
334 larger Catarrhini and the smaller Strepsirrhini and Platyrrhini, which both have larger caeca
335 compared to the Catarrhini (Fig. 12D). The difference in caecum size and shape between
336 these groups has been known for some time (Reider, 1936; Scott, 1980). One can only
337 speculate that for smaller species, the retention times necessary for allo-enzymatic fiber
338 digestion are more easily achieved by a larger dead-end structure such as the caecum rather
339 than by a larger colon.

340 Ultimately, primate clades have a typical mammalian macroscopic appearance of their
341 gastrointestinal tracts. The measure of intestinal length in primates matches broad statements
342 about differences between carnivorous and herbivorous species mentioned in the

343 **Introduction**, and, in the case of the large intestine, about the effect of natural diets on primate
 344 intestinal length. Within primates, however, dietary effects are found only across but not
 345 within clades, as models of the effect of diet on intestine length are only significant if the
 346 phylogenetic structure of the dataset is not accounted for. Within primate clades, dietary
 347 specialization **as measured in our dataset** has little power to explain intestinal length
 348 measures.

349

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 356 corresponding national and international protocols and requirements related to the use of
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358

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495 **Table 1** Average (\pm S.D.) body mass (g) and intestinal length (cm) measurements of primate species

| Species (Clade ^a) | Source | n | Body mass | Small intestine | Caecum | Colon | Large intestine ^b | Total intestine | %leaves in natural diet ^c | Diet quality index ^c |
|---------------------------------------|--------|----|------------------------|------------------|-----------------|------------------|------------------------------|------------------|---|------------------------------------|
| <i>Alouatta sara</i> (P) | 1 | 2 | 3950 (\pm 636) | 123 (\pm 11) | 6 (\pm 5) | 74 (\pm 3) | 80 (\pm 1) | 202 (\pm 10) | 50 | NA |
| <i>Alouatta seniculus</i> (P) | 2 | 1 | 635 | 239 | 12 | 58 | 70 | 309 | 48 | 328 |
| <i>Ateles paniscus</i> (P) | 1 | 1 | 8070 | 211 | 13 | 60 | 73 | 283 | 8 | 456 |
| <i>Callithrix argentata</i> (P) | 1 | 1 | 242 | 43 | 4 | 23 | 28 | 70 | 0 | NA |
| <i>Callithrix jacchus</i> (P) | 1,3 | 5 | 323 (\pm 30) | 58 (\pm 14) | 5 (\pm 1) | 25 (\pm 13) | 30 (\pm 12) | 88 (\pm 27) | 0 | 519 |
| <i>Cebuella pygmaea</i> (P) | 1 | 1 | 163 | 41 | 5 | 24 | 29 | 69 | 0 | 599 |
| <i>Cebus apella</i> (P) | 1 | 1 | 1750 | 145 | 5 | 34 | 39 | 184 | 16 | 543 |
| <i>Cheirogaleus medius</i> (S) | 1 | 2 | 197 (\pm 49) | 114 (\pm 0) | 5 (\pm 0) | 16 (\pm 3) | 21 (\pm 3) | 135 (\pm 3) | 4 | 549 |
| <i>Chiropotes satanas</i> (P) | 2 | 1 | 3130 | 282 | 19 | 48 | 67 | 349 | 1 | 504 |
| <i>Chlorocebus pygerythrus</i> (C) | 1 | 1 | 5300 | 176 | 7 | 67 | 74 | 250 | 10 | 526 |
| <i>Colobus guereza</i> (C) | 1 | 2 | 9750 (\pm 354) | 278 (\pm 57) | 8 (\pm 0) | 104 (\pm 22) | 112 (\pm 22) | 390 (\pm 79) | 61 | 348 |
| <i>Eulemur coronatus</i> (S) | 4 | 1 | 1580 | 99 | 14 | 55 | 69 | 168 | 3 | 493 |
| <i>Eulemur fulvus</i> (S) | 1 | 1 | 2500 | 91 | 16 | 57 | 73 | 164 | 25 | 396 |
| <i>Eulemur macaco</i> (S) | 1 | 2 | 1875 (\pm 530) | 125 (\pm 20) | 26 (\pm 6) | 66 (\pm 1) | 92 (\pm 5) | 217 (\pm 15) | 45 | 454 |
| <i>Gorilla gorilla</i> (C) | 1,5 | 2 | 154648 (\pm 102032) | 810 (\pm 299) | 18 ^d | 139 ^d | 162 (\pm 8) | 972 (\pm 307) | 61 | 359 |
| <i>Hapalemur griseus</i> (S) | 6 | 4 | 648 (\pm 106) | 86 (\pm 16) | 7 (\pm 4) | 50 (\pm 6) | 57 (N.R.) | 144 (N.R.) | 100 | 296 |
| <i>Homo sapiens</i> (C) | 9 | 6 | 65300 (\pm 14471) | 678 (\pm 138) | 8 (\pm 2) | 146 (\pm 16) | 154 (\pm 15) | 831 (\pm 144) | 30 | NA |
| <i>Lemur catta</i> (S) | 1 | 3 | 2117 (\pm 144) | 156 (\pm 43) | 24 (\pm 7) | 86 (\pm 24) | 111 (\pm 28) | 267 (\pm 71) | 32 | 432 |
| <i>Leontopithecus chrysomelas</i> (P) | 1 | 3 | 642 (\pm 98) | 101 (\pm 12) | 5 (\pm 1) | 27 (\pm 4) | 32 (\pm 4) | 133 (\pm 15) | 0 | 490 |
| <i>Macaca arctoides</i> (C) | 1 | 4 | 14525 (\pm 3495) | 297 (\pm 51) | 7 (\pm 2) | 125 (\pm 47) | 132 (\pm 45) | 429 (\pm 95) | 40 | NA |
| <i>Macaca cyclopis</i> (C) | 7 | 6 | 119667 (\pm 1783) | 230 (\pm 29) | 6 (\pm 1) | 79 (\pm 7) | 85 (\pm 7) | 315 (\pm 30) | 23 | 470 |
| <i>Macaca fuscata</i> (C) | 1 | 1 | 11970 | 434 | 5 | 121 | 126 | 560 | 32 | 372 |
| <i>Macaca nigra</i> (C) | 1 | 1 | 5600 | 295 | 8 | 113 | 121 | 415 | 2 | 596 |
| <i>Macaca sylvanus</i> (C) | 1 | 1 | 9625 | 334 | 7 | 163 | 170 | 504 | 11 | 430 |
| <i>Mandrillus sphinx</i> (C) | 1 | 1 | 23000 | 515 | 8 | 189 | 197 | 713 | 8 | 482 |
| <i>Mirza zaza</i> (S) | 1 | 1 | 336 | 82 | 4 | 34 | 38 | 120 | 38 | 420 |
| <i>Nomascus leucogenys</i> (C) | 1 | 1 | 6550 | 449 | 13 | 83 | 96 | 545 | 59 | 356 |
| <i>Pan paniscus</i> (C) | 1 | 1 | 37730 | 527 | 12 | 234 | 245 | 773 | 24 | 415 |
| <i>Papio hamadryas</i> (C) | 1 | 2 | 23250 (\pm 8132) | 426 (\pm 161) | 10 (\pm 2) | 148 (\pm 12) | 158 (\pm 10) | 585 (\pm 172) | 28 | 401 |
| <i>Pithecia pithecia</i> (P) | 1,2 | 3 | 1467 (\pm 339) | 191 (\pm 25) | 11 (\pm 4) | 56 (\pm 17) | 67 (\pm 21) | 258 (\pm 46) | 9 | 492 |
| <i>Pongo pygmaeus</i> (C) | 9 | 1 | 56250 | 559 | 23 | 328 | 351 | 910 | 48 | 458 |
| <i>Propithecus tattersalli</i> (S) | 6 | 1 | 2760 | 362 | 37 | 252 | 289 | 651 | 39 | 378 |
| <i>Propithecus verreauxi</i> (S) | 6 | 2 | 3890 (\pm 438) | 365 (\pm 38) | 39 (\pm 7) | 364 (\pm 42) | 403 (N.R.) | 767 (N.R.) | 53 | 302 |
| <i>Saguinus fuscicollis</i> (P) | 1 | 1 | 330 | 81 | 2 | 22 | 25 | 106 | 0 | 614 |
| <i>Saguinus oedipus</i> (P) | 1 | 4 | 460 (\pm 211) | 75 (\pm 8) | 5 (\pm 1) | 29 (\pm 9) | 33 (\pm 10) | 109 (\pm 17) | 0 | NA |
| <i>Saimiri boliviensis</i> (P) | 1 | 1 | 1003 | 144 | 5 | 18 | 22 | 166 | 0 | 500 |
| <i>Saimiri sciureus</i> (P) | 1,2,8 | 22 | 678 (\pm 117) | 104 (\pm 17) | 4 (\pm 1) | 12 (\pm 4) | 16 (\pm 4) | 120 (\pm 21) | 0 | 684 |
| <i>Symphalangus syndactylus</i> (C) | 1 | 1 | 8500 | 298 | 5 | 60 | 65 | 362.46 | 45 | 435 |
| <i>Theropithecus gelada</i> (C) | 1 | 1 | 11400 | 227 | 4 | 121 | 125 | 352 | 94 | 313 |
| <i>Trachypithecus vetulus</i> (C) | 1 | 1 | 5000 | 305 | 9 | 83 | 92 | 397 | 45 | 300 |
| <i>Varecia rubra</i> (S) | 1 | 1 | 4200 | 162 | 35 | 72 | 108 | 269 | 12 | 454 |
| <i>Varecia variegata</i> (S) | 1 | 1 | 3520 | 102 | 25 | 51 | 75 | 178 | 9 | 472 |

NA not available

497 ^a C Catarrhini, P Platyrrhini, S Strepsirrhini

498 ^bincluding caecum length

499 ^cfor sources, see Table S1

500 ^dcaecum & colon length only obtained from the present study

501 Sources: 1 present study; 2 Fooden (1964); 3 Caton, Hill, Hume and Crook (1996); 4 Schwitzer (2009); 5 Steiner (1954); 6 Campbell, Eisemann, Williams and Glenn (2000); 7 Makita et al. (1984); 8 Beischer and
502 Furry (1964); 9 Chivers (pers. comm.): for a previous study on mammal body composition (Navarrete et al. 2011), handwritten notes of Dr. Chivers were obtained of files used in preparation of his publication on
503 mammal digestive tract anatomy (Chivers and Hladik 1980); while these notes did not record the lengths of the intestines for the species included in that publication, they contained measurements of some additional
504 specimens.

505 **Table 2** Phylogenetic signal (λ) and allometric relationships between intestinal length
 506 measures (in cm) and body mass (BM, in kg) according to $\text{Length} = a \text{ BM}^b$ with 95%
 507 confidence intervals for parameter estimates from analyses using Phylogenetic Generalized
 508 Least Squares

| Length of | n | λ (95%CI) | a (95%CI) | b (95%CI) |
|----------------------|----|-------------------|-------------------|-------------------|
| Total intestine | 42 | 0.82 (0.47; 0.96) | 187 (123; 287) | 0.38 (0.30; 0.47) |
| <i>Strepsirrhini</i> | 11 | 1.00 (0.40; NA) | 217 (127; 371) | 0.47 (0.15; 0.79) |
| <i>Platyrrhini</i> | 14 | 0.72 (NA; NA) | 156 (134; 181) | 0.41 (0.30; 0.51) |
| <i>Catarrhini</i> | 17 | 0 (NA; 0.46) | 198 (146; 269) | 0.34 (0.24; 0.45) |
| Small intestine | 42 | 0.80 (0.37; 0.94) | 130 (86; 198) | 0.38 (0.29; 0.47) |
| <i>Strepsirrhini</i> | 11 | 1.00 (0.57; NA) | 141 (94; 213) | 0.31 (0.07; 0.56) |
| <i>Platyrrhini</i> | 14 | 0.66 (NA; 0.97) | 115 (93; 142) | 0.45 (0.30; 0.60) |
| <i>Catarrhini</i> | 17 | 0 (NA; 0.58) | 143 (101; 203) | 0.34 (0.22; 0.46) |
| Caecum | 42 | 0.92 (0.62; NA) | 7.1 (3.2; 15.9) | 0.41 (0.25; 0.56) |
| <i>Strepsirrhini</i> | 11 | 0 (NA; 0.79) | 12.3 (10.5; 14.5) | 0.76 (0.61; 0.91) |
| <i>Platyrrhini</i> | 14 | 0.77 (NA; NA) | 6.2 (4.5; 8.4) | 0.39 (0.18; 0.61) |
| <i>Catarrhini</i> | 17 | 0 (NA; 0.69) | 3.9 (2.1; 6.9) | 0.29 (0.09; 0.48) |
| Colon | 42 | 0.94 (0.76; 0.99) | 46 (23; 93) | 0.38 (0.25; 0.51) |
| <i>Strepsirrhini</i> | 11 | 1.00 (0.77; NA) | 61 (37; 100) | 0.60 (0.30; 0.89) |
| <i>Platyrrhini</i> | 14 | 0.75 (0.07; NA) | 31 (23; 42) | 0.29 (0.09; 0.49) |
| <i>Catarrhini</i> | 17 | 0 (NA; 0.81) | 49 (32; 76) | 0.33 (0.19; 0.48) |
| Large intestine | 42 | 0.94 (0.74; 1.00) | 55 (28; 106) | 0.38 (0.26; 0.50) |
| <i>Strepsirrhini</i> | 11 | 1.00 (0.60; NA) | 75 (47; 120) | 0.64 (0.35; 0.92) |
| <i>Platyrrhini</i> | 14 | 0.82 (0.20; NA) | 38 (29; 49) | 0.31 (0.13; 0.48) |
| <i>Catarrhini</i> | 17 | 0 (NA; 0.76) | 54 (36; 82) | 0.33 (0.19; 0.47) |

509 NA - no respective confidence limit available
 510 for GLS results, see Table S1



Eulemur fulvus

5 cm



Eulemur macaca

5 cm



Varecia rubra

5 cm



Varecia variegata

5 cm



Lemur catta

5 cm

511
512

Figure 1 Digestive tracts of various lemur species. Note the long caecum in all species.



Cheirogaleus medius

5 cm

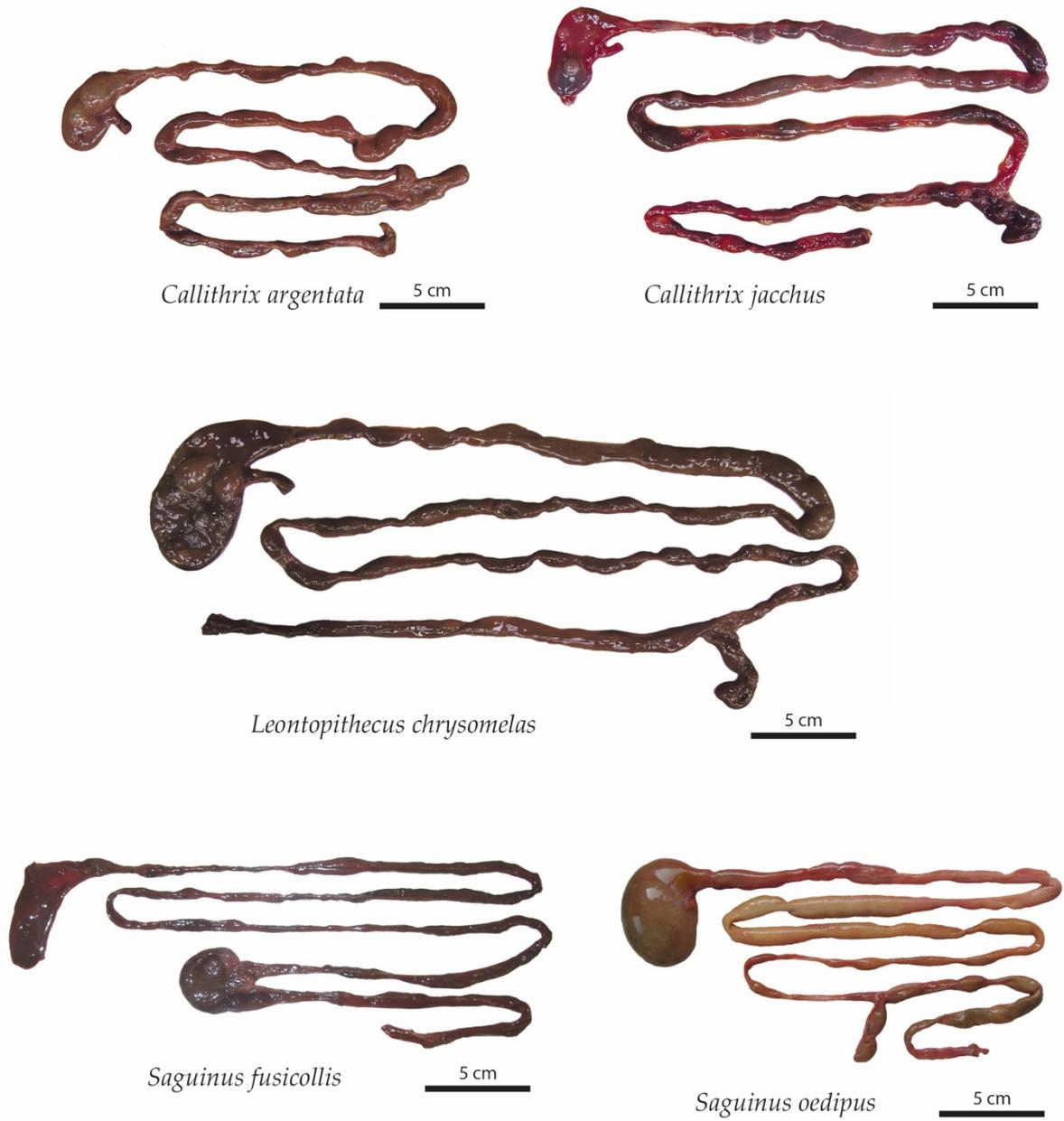


Mirza zaza

5 cm

513
514

Figure 2 Digestive tracts of two cheirogalid species.



515
516 **Figure 3** Digestive tracts of five callitrichid species.



Saimiri boliviensis

5 cm



Cebus apella

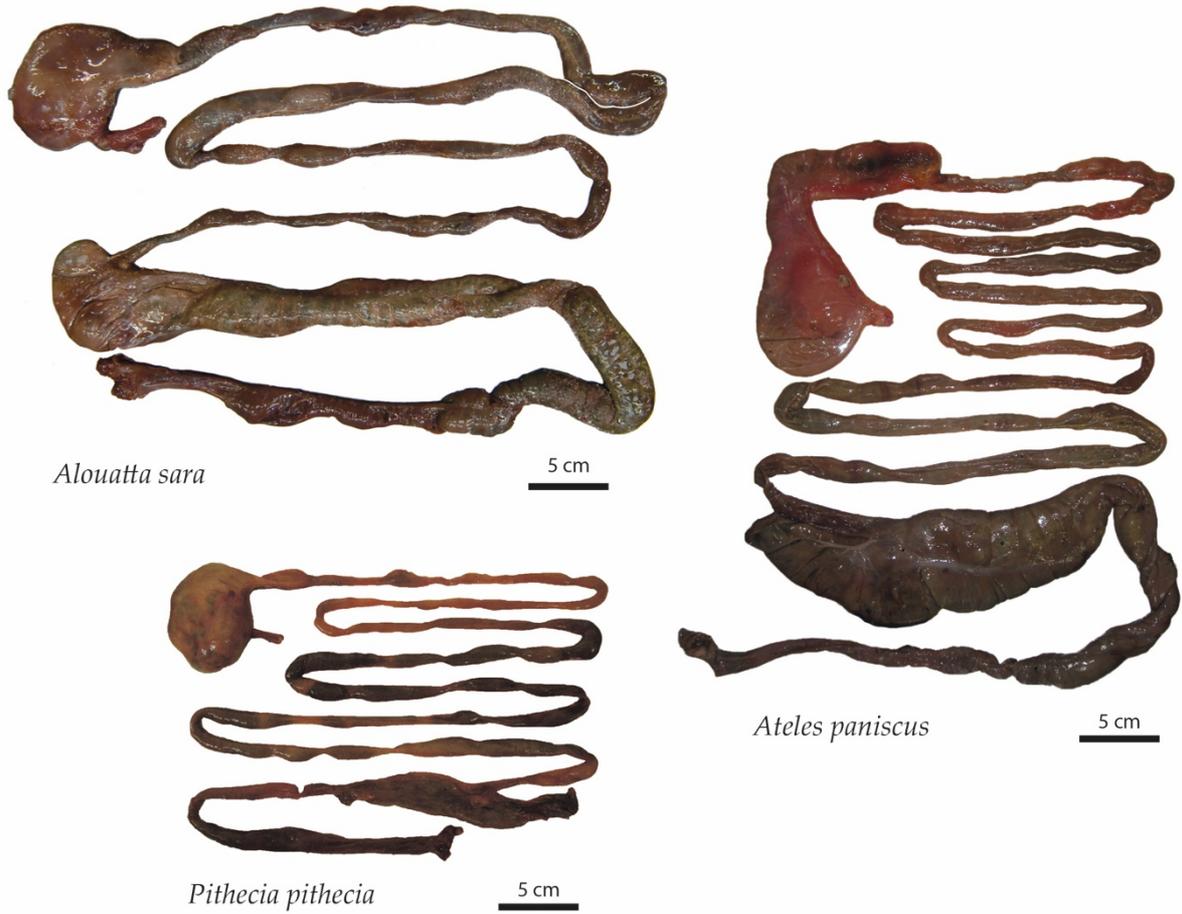
5 cm



Saimiri sciureus

5 cm

517
518 **Figure 4** Digestive tracts of three cebid species.



519
520 **Figure 5** Digestive tracts of two atelid and one pitheciid species.



Colobus guereza

5 cm



Trachypithecus vetulus

5 cm

521
522

Figure 6 Digestive tracts of two colobine species.



Chlorocebus pygerythrus

5 cm

523
524 **Figure 7** Digestive tract of *Chlorocebus*.



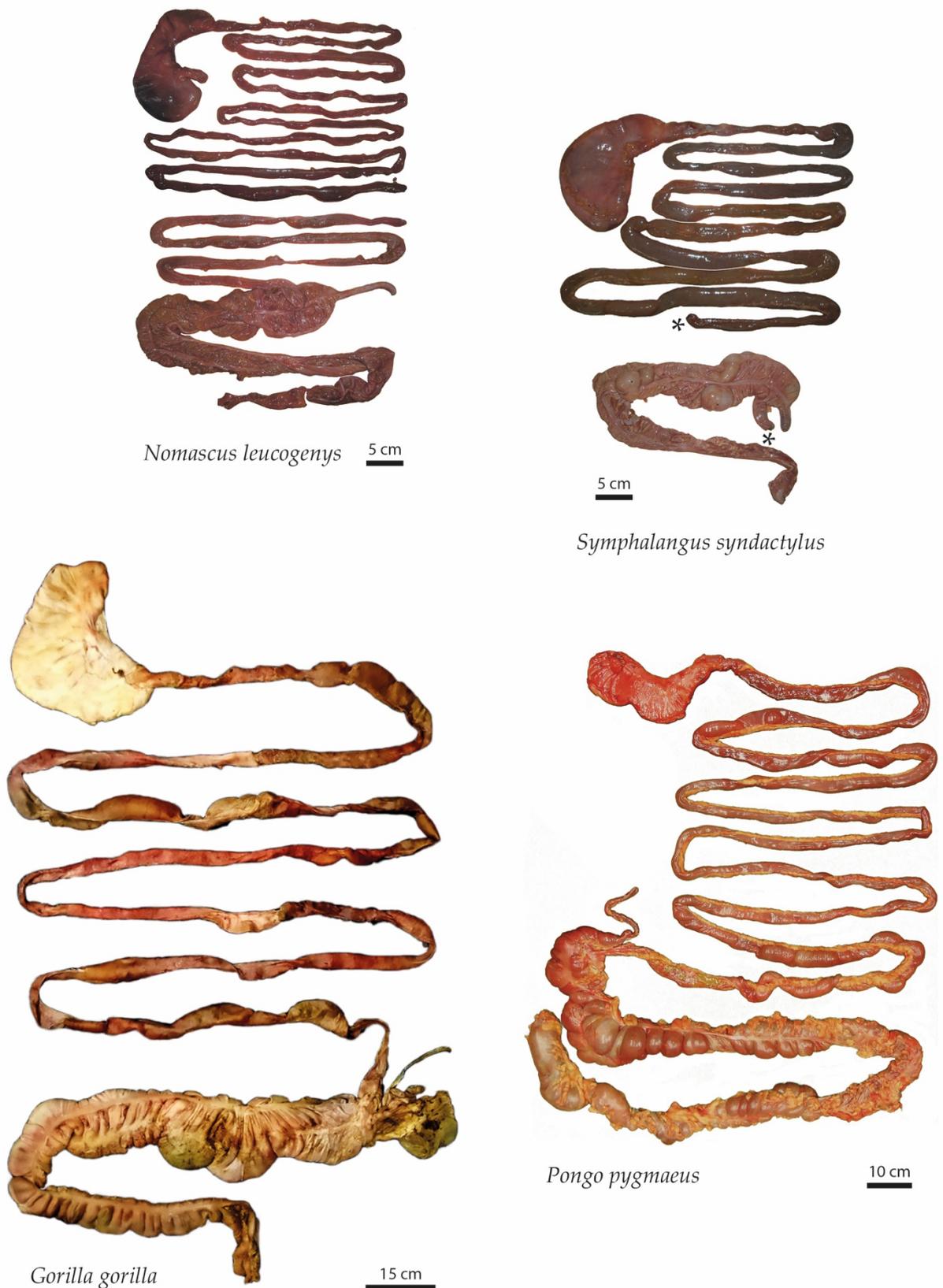
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Figure 8 Digestive tracts of four macaque species. The * indicates where the small intestine was linked before dissection.



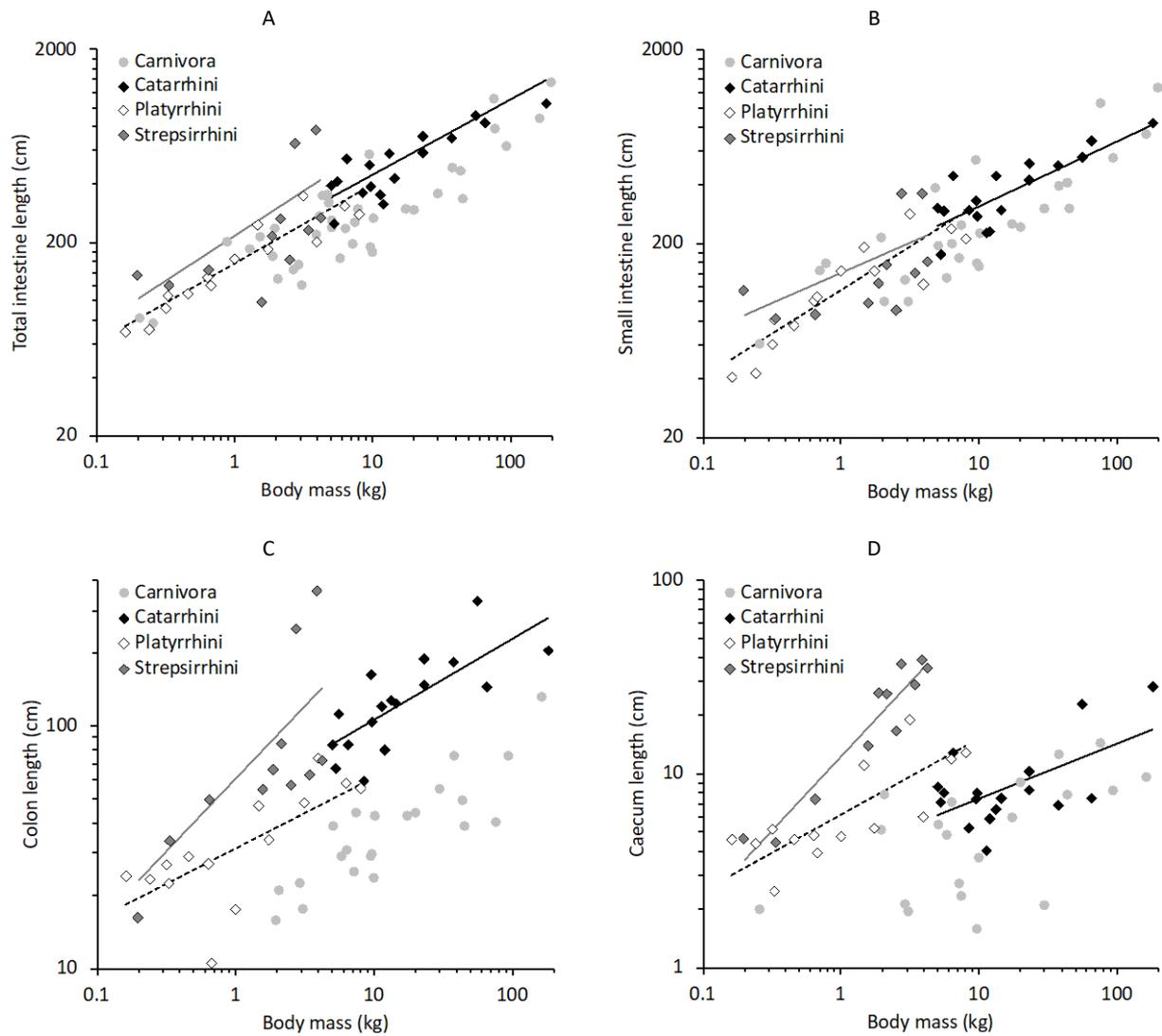
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Figure 9 Digestive tracts of three baboon species. The * marks where the small intestine was linked before dissection.



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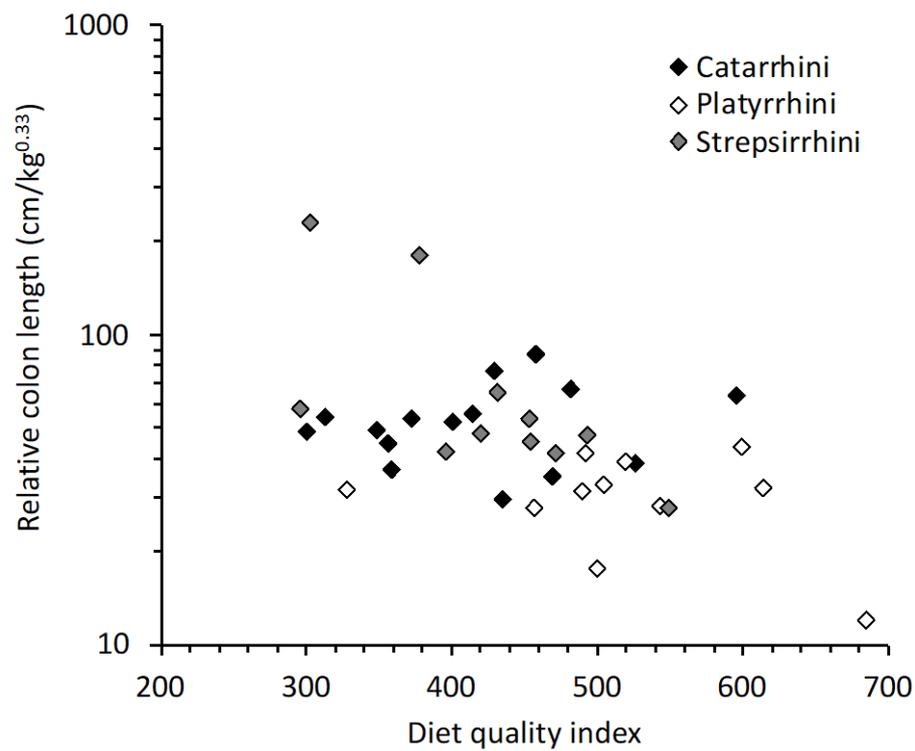
Figure 10 Digestive tracts in four ape species. The * marks where the small intestine was linked before dissection. Note the caecal appendix in all four species.



535

536 **Figure 11** Relationship between body mass and A) total intestinal length, B) small intestinal
 537 length, C) colon length and D) caecum length in primates as compared to terrestrial
 538 carnivorans (from McGrosky et al., 2016). Regression lines from Phylogenetic Generalized
 539 Least Squares (Table 2). Straight black line = Catarrhini; black dotted line = Platyrrhini; grey
 540 line = Strepsirrhini. The two outlier lemurs in (A-C) are sifakas (*Propithecus* spp.).

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543 **Figure 12** Relationship between the diet quality index and the relative length of the colon in
 544 species of three different primate clades. Note that while there is a negative relationship in
 545 the overall data, this is not the case in the individual clades, leading to a non-significant result
 546 when accounting for the phylogenetic structure of the data (see Results). The two outliers with
 547 a high relative colon length in the Strepsirrhini are from one genus (*Propithecus* spp.); the
 548 two outliers in the Platyrrhini with a low relative colon length are also from one genus
 549 (*Saimiri* spp.).

550 **Supporting Information**

551 **Gross intestinal morphometry and allometry in primates**

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554 **Table S1** Sources for diet information

555

556 ACHTUNG: alles normal ist aus den Van Woerden Papers bzw. aus der Excel-Tabelle; **gelb = keine Ahnung, woher (incl. der Angabe im**
 557 **Excel); blau = Angabe aus Excel, obwohl Quelle in Van Woerden Papers; pink = all the worlds primates – kann man da die direkte quelle**
 558 **rausziehen?**

| Species (Clade ^a) | Source |
|---------------------------------------|--|
| <i>Alouatta sara</i> (P) | Neville, M. K., Glander, K. E., Braza, F., & Rylands, A. B. (1988). The howling monkeys, genus <i>Alouatta</i> . In <i>Ecology and behavior of neotropical primates, vol 2</i> (ed. R. A. Mittermeier, A. B. Rylands, A. Coimbra-Filho, G. A. B. Fonseca). Washington DC: World Wildlife Fund. |
| <i>Alouatta seniculus</i> (P) | Julliot, C., & Sabatier, D. (1993). Diet of the red howler monkey (<i>Alouatta seniculus</i>) in French-Guiana. <i>Int J Primatol</i> 14:527-550. Di Fiore, A., & Campbell, C. J. (2007). The atelines: variation in ecology, behavior, and social organisation. In <i>Primates in perspective</i> (ed. C. J. Campbell, A. Fuentes, K. C. McKinnon, M. Panger, S. K. Bearder). New York: Oxford University Press. |
| <i>Ateles paniscus</i> (P) | van Roosmalen, M. G. M. (1980). <i>Habitat preferences, diet, feeding strategy and social organization of the black spider monkey (Ateles paniscus paniscus Linnaeus 1758) in Surinam</i> . PhD thesis. Landbouwhogeschool Wageningen: Leersum. Di Fiore, A., & Campbell, C. J. (2007). The atelines: variation in ecology, behavior, and social organisation. In <i>Primates in perspective</i> (ed. C. J. Campbell, A. Fuentes, K. C. McKinnon, M. Panger, S. K. Bearder). New York: Oxford University Press. |
| <i>Callithrix argentata</i> (P) | xxx (Willems and van Schaik 2015?) |
| <i>Callithrix jacchus</i> (P) | de Castro, C. S. S., & Araújo, A. (2007). Diet and feeding behavior of marmoset, <i>Callithrix jacchus</i> . <i>Brazilian Journal of Ecology</i> , 7, 14-19. Digby, L. J., Ferrari, S. F., & Saltzman, W. (2007). Callitrichines: the role of competition in cooperatively breeding species. In <i>Primates in perspective</i> (ed. C. J. Campbell, A. Fuentes, K. C. McKinnon, M. Panger, S. K. Bearder). New York: Oxford University Press. |
| <i>Cebuella pygmaea</i> (P) | xxx (Van Woerden 2010?) |
| <i>Cebus apella</i> (P) | Zhang, S. Y. (1995). Activity and ranging patterns in relation to fruit utilization by brown capuchins (<i>Cebus apella</i>) in French-Guiana. <i>International Journal of Primatology</i> , 16, 489-507. |
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| <i>Chiropotes satanas</i> (P) | Norconk, M. A. (1996). Seasonal variation in the diets of white-faced and bearded sakis (<i>Pithecia pithecia</i> and <i>Chiropotes satanas</i>) in Guri Lake, Venezuela. In <i>Adaptive radiations of neotropical primates</i> (ed. M. A. Norconk, A. L. Rosenberger, P. A. Garber). New York: Plenum Press. |
| <i>Chlorocebus pygerythrus</i> (C) | Enstam, K. L., & Isbell, L. (2007). The guenons (<i>genus Cercopithecus</i>) and their allies: behavioral ecology of polyspecific associations. In <i>Primates in perspective</i> (ed. C. J. Campbell, A. Fuentes, K. C. McKinnon, M. Panger, S. K. Bearder). New York: Oxford University Press. |
| <i>Colobus guereza</i> (C) | Plumptre, A. J. (2006). The diets, preferences, and overlap of the primate community in the Budongo Forest Reserve, Uganda. In <i>Primates of western Uganda</i> (ed. N. E. Newton-Fisher, H. Notman, J. D. Paterson & V. Reynolds). New York: Springer; Fashing, P. J. (2001). Feeding ecology of guerezas in the Kakamega Forest, Kenya: The importance of <i>Moraceae</i> fruit in their diet. <i>International Journal of Primatology</i> , 22, 579-609. |
| <i>Eulemur coronatus</i> (S) | Rowe N., & Myers M. (2011). <i>All the world's primates</i> . Rhode Island: Primate Conservation Inc. |
| <i>Eulemur fulvus</i> (S) | Rasmussen, M. A. (1999). <i>Ecological influences on activity cycle in two cathemeral primates, the mongoose lemur (Eulemur mongoz) and the common brown lemur (Eulemur fulvus fulvus)</i> . PhD diss. Duke University, Durham, NC; Tarnaud, L. (2006). Feeding behavior of lactating brown lemur females (<i>Eulemur fulvus</i>) in Mayotte: influence of infant age and plant phenology. <i>American Journal of Primatology</i> , 68, 966-977. |
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| <i>Homo sapiens</i> (C) | Wilman H., Belmaker J., Simpson J., de la Rosa C., Rivadeneira M. M., & Jetz W. (2014). EltonTraits 1.0: Species-level foraging attributes of the world's birds and mammals. <i>Ecology</i> , 95, 2027-2027. |
| <i>Lemur catta</i> (S) | Rowe N., & Myers M. (2011). <i>All the world's primates</i> . Rhode Island: Primate Conservation Inc. |
| <i>Leontopithecus chrysomelas</i> (P) | xxx (Willems and van Schaik 2015?) |

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560 **Table S2** Allometric relationships between intestinal length measures (in cm) and body mass
 561 (BM, in kg) according to $\text{Length} = a \text{ BM}^b$ with 95% confidence intervals for parameter
 562 estimates from analyses using Generalized Least Squares (i.e., not accounting for phylogeny)

| Length of | n | a (95%CI) | b (95%CI) |
|----------------------|----|-------------------|-------------------|
| Total intestine | 42 | 168 (148; 190) | 0.40 (0.34; 0.46) |
| <i>Strepsirrhini</i> | 11 | 188 (133; 367) | 0.40 (0.08; 0.73) |
| <i>Platyrrhini</i> | 14 | 152 (137; 168) | 0.40 (0.32; 0.49) |
| <i>Catarrhini</i> | 17 | 198 (146; 269) | 0.34 (0.24; 0.45) |
| Small intestine | 42 | 117 (103; 132) | 0.41 (0.35; 0.46) |
| <i>Strepsirrhini</i> | 11 | 125 (93; 167) | 0.29 (0.01; 0.56) |
| <i>Platyrrhini</i> | 14 | 111 (96; 128) | 0.44 (0.32; 0.56) |
| <i>Catarrhini</i> | 17 | 143 (101; 203) | 0.34 (0.22; 0.46) |
| Caecum | 42 | 7.5 (5.7; 9.7) | 0.16 (0.04; 0.29) |
| <i>Strepsirrhini</i> | 11 | 12.3 (10.5; 14.5) | 0.76 (0.61; 0.91) |
| <i>Platyrrhini</i> | 14 | 6.1 (5.0; 7.4) | 0.34 (0.18; 0.51) |
| <i>Catarrhini</i> | 17 | 3.9 (2.1; 6.9) | 0.29 (0.09; 0.48) |
| Colon | 42 | 41 (34; 49) | 0.41 (0.32; 0.50) |
| <i>Strepsirrhini</i> | 11 | 54 (37; 79) | 0.62 (0.26; 0.98) |
| <i>Platyrrhini</i> | 14 | 31 (25; 38) | 0.31 (0.14; 0.47) |
| <i>Catarrhini</i> | 17 | 49 (32; 76) | 0.33 (0.19; 0.48) |
| Large intestine | 42 | 49 (41; 59) | 0.38 (0.29; 0.47) |
| <i>Strepsirrhini</i> | 11 | 69 (48; 98) | 0.66 (0.34; 0.98) |
| <i>Platyrrhini</i> | 14 | 38 (31; 45) | 0.32 (0.17; 0.47) |
| <i>Catarrhini</i> | 17 | 54 (36; 82) | 0.33 (0.19; 0.47) |

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