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Year: 2013

Assessing the Jarman–Bell Principle: Scaling of intake, digestibility, retention time and gut fill with body mass in mammalian herbivores

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Abstract: Differences in allometric scaling of physiological characters have the appeal to explain species diversification and niche differentiation along a body mass (BM) gradient - because they lead to different combinations of physiological properties, and thus may facilitate different adaptive strategies. An important argument in physiological ecology is built on the allometries of gut fill (assumed to scale to BM1.0) and energy requirements/intake (assumed to scale to BM0.75) in mammalian herbivores. From the difference in exponents, it has been postulated that the mean retention time (MRT) of digesta should scale to BM1.0-0.75 = BM0.25. This has been used to argue that larger animals have an advantage in digestive efficiency and hence can tolerate lower-quality diets. However, empirical data does not support the BM0.25 scaling of MRT, and the deduction of MRT scaling implies, according to physical principles, no scaling of digestibility; basing assumptions on digestive efficiency on the thus-derived MRT scaling amounts to circular reasoning. An alternative explanation considers a higher scaling exponent for food intake than for metabolism, allowing larger animals to eat more of a lower quality food without having to increase digestive efficiency; to date, this concept has only been explored in ruminants. Here, using data for 77 species in which intake, digestibility and MRT were measured (allowing the calculation of the dry matter gut contents DMC), we show that the unexpected shallow scaling of MRT is common in herbivores and may result from deviations of other scaling exponents from expectations. Notably, DMC have a lower scaling exponent than 1.0, and the 95% confidence intervals of the scaling exponents for intake and DMC generally overlap. Differences in the scaling of wet gut contents and dry matter gut contents confirm a previous finding that the dry matter concentration of gut contents decreases with body mass, possibly compensating for the less favourable volume-surface ratio in the guts of larger organisms. These findings suggest that traditional explanations for herbivore niche differentiation along a BM gradient should not be based on allometries of digestive physiology. In contrast, they support the recent interpretation that larger species can tolerate lower-quality diets because their intake has a higher allometric scaling than their basal metabolism, allowing them to eat relatively more of a lower quality food without having to increase digestive efficiency.

DOI: https://doi.org/10.1016/j.cbpa.2012.09.018

Posted at the Zurich Open Repository and Archive, University of Zurich ZORA URL: https://doi.org/10.5167/uzh-67071 Journal Article Accepted Version

Originally published at:

Müller, Dennis W H; Codron, Daryl; Meloro, Carlo; Munn, Adam; Schwarm, Angela; Hummel, Jürgen; Clauss, Marcus (2013). Assessing the Jarman–Bell Principle: Scaling of intake, digestibility, retention time and gut fill with body mass in mammalian herbivores. Comparative Biochemistry and Physiology Part A: Molecular Integrative Physiology, 164(1):129-140. DOI: https://doi.org/10.1016/j.cbpa.2012.09.018

1	Assessing the Jarman-Bell Principle: scaling of intake, digestibility, retention time and
2	gut fill with body mass in mammalian herbivores
3	
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24	Running head: Herbivore digestive allometry
25	

26 Abstract

27 Differences in allometric scaling of physiological characters have the appeal to explain 28 species diversification and niche differentiation along a body mass (BM) gradient - because 29 they lead to different combinations of physiological properties, and thus may facilitate 30 different adaptive strategies. An important argument in physiological ecology is built on the allometries of gut fill (assumed to scale to BM^{1.0}) and energy requirements/intake (assumed to 31 scale to BM^{0.75}) in mammalian herbivores. From the difference in exponents, it has been 32 postulated that the mean retention time (MRT) of digesta should scale to $BM^{1.0-0.75} = BM^{0.25}$. 33 This has been used to argue that larger animals have an advantage in digestive efficiency and 34 hence can tolerate lower-quality diets. However, empirical data does not support the BM^{0.25} 35 scaling of MRT, and the deduction of MRT scaling implies, according to physical principles, 36 37 no scaling of digestibility; basing assumptions on digestive efficiency on the thus-derived MRT scaling amounts to circular reasoning. An alternative explanation considers a higher 38 39 scaling exponent for food intake than for metabolism, allowing larger animals to eat more of a 40 lower quality food without having to increase digestive efficiency; to date, this concept has 41 only been explored in ruminants. Here, using data for 77 species in which intake, digestibility 42 and MRT were measured (allowing the calculation of the dry matter gut contents DMC), we show that the unexpected shallow scaling of MRT is common in herbivores and may result 43 44 from deviations of other scaling exponents from expectations. Notably, DMC have a lower 45 scaling exponent than 1.0, and the 95% confidence intervals of the scaling exponents for 46 intake and DMC generally overlap. Differences in the scaling of wet gut contents and dry 47 matter gut contents confirm a previous finding that the dry matter concentration of gut 48 contents decreases with body mass, possibly compensating for the less favourable volume-49 surface ratio in the guts of larger organisms. These findings suggest that traditional 50 explanations for herbivore niche differentiation along a BM gradient should not be based on 51 allometries of digestive physiology. In contrast, they support the recent interpretation that 52 larger species can tolerate lower-quality diets because their intake has a higher allometric 53 scaling than their basal metabolism, allowing them to eat relatively more of a lower quality 54 food without having to increase digestive efficiency.

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56 Key words: Jarman-Bell Principle; herbivory; body size; diet quality; digestion; allometry;

57 feeding ecology; digestive physiology

59 Introduction

60 The Jarman-Bell Principle

61 The scaling relationships of digestive features with body mass (BM) in herbivores are a 62 fundamental part of our interpretation of herbivore feeding ecology, interspecific niche 63 differentiation and intraspecific niche segregation due to sexual dimorphism. Geist (1974) 64 explained how two PhD studies, which resulted in two publications (Bell, 1971; Jarman, 65 1974), crystallized an understanding of the ways in which BM affects the ecology of ungulates of the African savannahs. These studies have become the springboard for most 66 67 analyses of mammal herbivore ecological interactions that include a range of BM (du Toit, 2005). The so-called 'Jarman-Bell Principle' (JBP) (Geist, 1974) has been applied to other 68 69 animal groups such as primates (Gaulin, 1979), fruit bats (Fleming, 1991) or baleen whales 70 (Tershy, 1992), and predicts that BM constrains nutritional ecology – in terms of the diet 71 quality that larger animals can or must tolerate, as well as their digestive efficiency. That 72 larger animals commonly ingest diets of lower quality has been demonstrated repeatedly (e.g. 73 Owen-Smith, 1988; Codron et al., 2007), and that they often must tolerate poor-quality foods 74 is usually not an issue of debate. This is considered a consequence of large animals' higher 75 absolute food requirements that prevent them from foraging selectively in terms of time 76 constraints, and also a consequence of larger, more 'clumsy' feeding apparatus that again 77 prevents them from foraging selectively in terms of picking out individual food items or plant 78 parts (Owen-Smith, 1988; Shipley, 2007).

79

80 A fact often used to explain the increased dietary tolerance (measured as the range in fibre 81 and/or protein content of the diet the animal can tolerate) of larger species are the decreased 82 mass-specific metabolic demands - because the daily requirements for energy (Nagy et al., 83 1999), nitrogen (Robbins, 1993) and even minerals (Rucker and Storms, 2002) scale to metabolic body mass (BM^{0.75}). This scaling effect means that small-bodied species require 84 85 more energy and nutrients per day and per unit of BM than do large-bodied forms (Geist, 86 1974). However, without information about the scaling of energy/nutrient intake, knowledge 87 about the scaling of requirements allows no further insight. Concepts about niche 88 diversification along a BM gradient must make an assumption about the difference in scaling 89 between requirement and intake. Given that larger animals ingest lower-quality diets, a 90 similar scaling of intake and requirements would make it necessary that digestive efficiency 91 has a positive scaling, i.e. it should increase with BM. This could be achieved either by 92 selecting a more digestible diet (which violates the assumption of decreasing diet quality), or

by measures of digestive physiology that increase digestive efficiency, such as prolonged
digesta retention. In contrast, if we postulate that there is no scaling of digestive efficiency

95 with BM, then intake would have to scale to BM at a higher exponent than requirements, in

- 96 order to compensate for the lower diet quality in larger animals. Historically, the JBP has
- 97 more often been linked to variation in digestive efficiency than to variation in intake levels.
- 98

99 When developing this approach of the JBP, food intake was, without empirical testing, 100 assumed to scale to body mass in a similar way as animal metabolism (explicitly in Demment, 101 1983; implicitly in Demment and Van Soest, 1985; note that Van Soest, 1994 does not make 102 an explicit assumption about intake scaling but rests his argument on the scaling of gut fill 103 only). In large datasets, food intake (measured as dry matter intake, DMI, or as gross energy intake) was actually often found to scale to BM^{0.75} across a wide range of mammalian species 104 105 (reviewed in Clauss et al., 2007a; Meyer et al., 2010), and thus resembles the scaling of 106 metabolism and requirements. Therefore, the approach that focuses on scaling effects on 107 digestive physiology appears as the logical next step. Including gut fill in these considerations 108 has extended the JBP to specifically comprise not only metabolic scaling, but also digestive 109 physiology. Gut fill (measured as wet matter content, WMC) or dry matter content (DMC) 110 has been suggested to be a constant fraction of BM in herbivores (Parra, 1978; Demment and 111 Van Soest, 1985; Illius and Gordon, 1992; Justice and Smith, 1992; Clauss et al., 2007a). If gut fill actually scales to BM^{1.00}, and food intake scales to BM^{0.75}, these different scaling 112 113 effects result in a larger gut fill per unit food intake with increasing BM. This should in theory 114 lead to an increase in the time digesta is retained in the gastrointestinal tract (measured as 115 mean retention time, MRT) with increasing BM (Demment and Van Soest, 1985; Illius and Gordon, 1992; Robbins, 1993; McNab, 2002). Explicitly, it is thought that MRT should scale 116 to $BM^{(1.00-0.75)} = BM^{0.25}$. Because MRT is positively linked to the digestive efficiency of a 117 herbivore (Foose, 1982; Udén and Van Soest, 1982; Clauss et al., 2007b), this concept has 118 119 been invoked to postulate that larger herbivores can use food of a lower quality due to a 120 hypothetical capacity for more efficient digestion (Demment and Van Soest, 1985; Illius and 121 Gordon, 1992; Gordon and Illius, 1996). In this way, the JBP delivers an elegant explanation 122 (higher digestive efficiency in larger animals) for a common observation (lower diet quality in 123 larger animals), and represents what may be the most prominent set of allometric 124 considerations in large animal ecology, including theoretical approaches to understanding 125 intraspecific size dimorphism (Short, 1963; Prins and Geelen, 1971; Sinclair, 1977; Hanley

126	and Hanley, 1980; Hanley, 1982; Barboza and Bowyer, 2000; du Toit, 2005; Shannon et	al.,
127	2006; Woolley et al., 2011).	
128		
129	Empirical evidence for the physiological predictions of the JBP, however, is scarce. While	e the
130	findings that wet gut contents scale to BM ^{1.00} and food intake scales to BM ^{0.75} are mostly	
131	undisputed, digesta retention has repeatedly been shown, in various datasets, not to scale	to
132	BM ^{0.25} (Clauss et al., 2007a; Clauss et al., 2009; Franz et al., 2011a; Müller et al., 2011;	
133	Steuer et al., 2011), with the exception of mammalian caecum fermenters (Clauss et al.,	
134	2007a, but see the results of phylogenetically informed statistics in this study down below	v).
135	Additionally, there is little indication for an increase in digestive efficiency with BM (Just	tice
136	and Smith, 1992; Wenninger and Shipley, 2000; Pérez-Barbería et al., 2004; Clauss et al.	,
137	2009; Steuer et al., 2012). These findings represent a serious conceptual problem for	
138	herbivore digestive physiology, because the link between gut fill, food intake and digesta	
139	retention is mathematically sound.	
140		
141	Linking digesta retention to gut capacity, food intake and digestibility	
142	Holleman & White (1989) provided the underlying equation to this problem, based on the	•
143	'Stewart-Hamilton Principle' or the 'Occupancy Principle' (Steele, 1971; Shipley and Cla	ark,
144	1972), where	
145	indDMC [kg] = faecal output [kg/h] * MRT _{particle} GIT [h]	1)
146	with indDMC being the indigestible (or undigested) DM content of the whole gastrointes	tinal
147	tract (GIT), and MRT _{particle} GIT the mean retention time of an indigestible particle marker	in
148	the whole GIT. Because faecal output is a function of food intake (measured as dry matte	r
149	intake DMI) and apparent digestibility of dry matter (aD DM) or, in other terms, DMI an	d the
150	apparent indigestibility of DM (aID DM), we get	
151	faecal output $[kg/h] = DMI [kg/h] * (aID DM)$	2)
152	and thus gut fill, food intake and retention time are closely linked as	
153	indDMC [kg] = DMI [kg/h] * aID DM * MRT _{particle} GIT [h] $($	3)
154	Solving this equation for MRT yields	
155	$MRT_{particle}GIT [h] = indDMC [kg] / (DMI [kg/h] * aID DM) $	4)
156	Assuming scaling laws for indDMC, DMI and aID DM with body mass with the exponen	ıts a,
157	b and c, respectively, it follows that	
158	$MRT_{particle}GIT \sim BM^{a} / (BM^{b} * BM^{c}) \text{ or } MRT_{particle}GIT \sim BM^{a-(b+c)} $	5)

- 159 Under the assumption that indDMC scales as does wet matter contents, i.e. to BM^{1.00}, and
- 160 DMI scales to $BM^{0.75}$, we can thus assume that $MRT_{particle}GIT$ should scale to $BM^{0.25}$ if
- 161 *digestive efficiency does not change with BM*. Note that an increase in digestibility with

162 increasing BM (i.e., a decrease in indigestibility, or a negative exponent c) would lead to a

- 163 higher scaling exponent for MRT_{particle}GIT, and a decrease in digestibility with increasing BM
- 164 would lead to a lower scaling exponent for MRT_{particle}GIT. This is because a lower
- 165 digestibility would result in more material in the GIT that is pushed along due to the
- 166 continuous food intake, and hence make any particular portion of the indigestible digesta be
- 167 excreted relatively sooner (Hummel and Clauss, 2011).
- 168

169 The calculation of gut fill can be expanded to include not only indigestible gut fill, but also

- total gut fill (dry matter contents DMC) (Holleman and White, 1989). For that, the digestible
- 171 portion of DMC (dDMC) has to be added to indDMC. To do so, one has to make an
- assumption regarding the course of digestion. If one assumes digestion to be exponential over
- 173 MRT (with most of digestion occurring at the beginning of MRT and slowly 'fading out'),
- then the estimated dDMC in the total GIT will be less than if one assumes digestion to be
- 175 linear across MRT. Holleman and White (1989) and Munn et al. (2012) showed that assuming
- a linear course of digestion across MRT yields better results when validated against empirical
- 177 measures in slaughtered animals, albeit using small sample sizes. Because assuming a linear
- 178 course of digestion across MRT means that dDMC is assumed to be present in the GIT during
- 179 half of MRT, it follows that

180
$$dDMC [kg] = DMI [kg/h] * aD DM * (MRT_{particle}GIT [h] / 2)$$
 (6).

- 181 Combining equation (3) and (6) to calculate total DMC yields,
- 182 DMC [kg] = DMI [kg/h] * MRT_{particle}GIT [h] * (aID DM + 0.5 aD DM) (7) 183 Solving the equation for MRT_{particle}GIT [h] yields
- 184 $MRT_{particle}GIT[h] = DMC[kg] / DMI[kg/h] * (1 / (aID DM + 0.5 aD DM))$ (8)
- Assuming scaling laws for DMC, DMI and aD DM with body mass with the exponents d, band c, respectively, it follows that

187
$$MRT_{particle}GIT \sim BM^{d} / BM^{b} * (1 / (BM^{-c} + 0.5 BM^{c})) \text{ or } MRT_{particle}GIT \sim BM^{d-b+c}$$
188 (9)

- 189 Assuming that DMC scales similar as wet matter contents, i.e. to BM^{1.00}, and DMI scales to
- 190 $BM^{0.75}$, we can thus again assume that MRT_{particle}GIT should scale to $BM^{0.25}$ *if digestive*
- 191 efficiency does not change with BM. Note that again, an increase in digestibility with BM
- 192 (i.e., a positive c) would lead to a higher scaling exponent for MRT_{particle}GIT, and a decrease

in digestibility with BM would lead to a lower scaling exponent for MRT_{particle}GIT. One
important conclusion from these derivations is that if one uses the scaling of intake and gut
fill to make any conclusions about digesta retention, one must not use the result to make
inferences about digestive efficiency - because an assumption about the scaling of
digestibility is made already (explicitly or implicitly) to derive the scaling of digesta
retention. Deriving a scaling for digesta retention in this way and then using it to explain a
pattern of digestive efficiency with body mass amounts to circular reasoning.

200

201 Re-assessing data from herbivore feeding trials

202 Given these considerations, a discrepancy in the scaling of gut contents, food intake, and 203 digesta retention, as indicated by empirical data, warrants a re-assessment of both empirical 204 datasets and, potentially, a revision of our understanding of herbivore digestive allometry. An 205 intuitive reason for this discrepancy might be that whereas it is common practice to measure 206 both food intake and digesta retention in the same experiment, gut fill is usually not measured 207 simultaneously, as most experiments do not include the dissection of the experimental 208 animals. The discrepancy of the scaling exponents therefore might simply result from the fact 209 that different datasets have been used to generate these allometries (Clauss et al., 2007a); 210 nevertheless, the discrepancy requires further explanation. If in addition to measures of food 211 intake and digesta retention, the apparent digestibility of dry matter is also measured in the 212 same feeding trial, the derivation of indigestible and complete gut fill (as DMC) as indicated 213 in the equations above is possible. Although this has been done repeatedly in individual 214 studies to compare selected species, or make intraspecific or inter-clade comparisons (Baker 215 and Hobbs, 1987; Gross et al., 1996; Behrend et al., 2004; Munn and Dawson, 2006; Munn 216 and Barboza, 2008; Schwarm et al., 2009a; Clauss et al., 2010b; Franz et al., 2011b; Sawada 217 et al., 2011; Fritz et al., 2012; Munn et al., 2012), a large-scale comparison of mammalian 218 herbivore gut fill has not been performed so far. Therefore, it is the aim of this study to 219 explore the scaling of food intake, digesta retention, apparent digestibility and the calculated 220 gut fill in mammalian herbivores, using available literature data. Because these variables are 221 used to calculate gut fill according to the principles outlined above, we expect that their 222 scaling relationships also follow these principles.

223

224 The following hypotheses guided our investigation:

- (i) Across the whole mammal body size range, food intake measured as dry matter
 intake (DMI) has a scaling exponent close to BM^{0.75} (reviewed in Clauss et al.,
 2007a).
- 228 Apparent digestibility generally does not scale with body mass (i.e. to $BM^{0.00}$) (ii) 229 (Justice and Smith, 1992; Wenninger and Shipley, 2000; Pérez-Barbería et al., 230 2004; Clauss et al., 2009; Steuer et al., 2012), and therefore the scaling exponents of gut fill (BM^a), food intake (BM^b) and digesta retention (BM^d) are associated 231 according to d = a-b. It should be noted, however, that any conclusions concerning 232 233 the apparent digestibility must be considered in the context of the food sources that 234 were used across experiments. When collating data for a large variety of species 235 from the literature, as we do here, it is highly likely that data from smaller species, 236 which are often kept on artificial feeds for experimental purposes, is combined 237 with data from larger species, which are more often kept on roughage-only diets 238 that are generally less digestible. Thus, a putative negative allometry of 239 digestibility (a decrease with body mass) would be an indication of this uneven 240 distribution of diet quality. How close this potential decrease in diet quality with 241 body size in the dataset actually might resemble the general decrease of diet 242 quality with body size that is supposed to occur in the wild is difficult to judge. 243 Therefore, if scaling is detected between body mass and digestibility, then the scaling exponents of gut fill (BM^a), food intake (BM^b), digestibility (BM^c) and 244 digesta retention (BM^d) are associated according to d = a-b+c. 245
- (iii) The low empirical overall scaling exponent for digesta retention (Clauss et al.,
 2007a; Clauss et al., 2009; Müller et al., 2011; Steuer et al., 2011) results from the
 fact that gut fill, measured as dry matter contents (DMC), does *not* scale to BM^{1.00}
 but to a lower exponent.
- 250 (iv) In parallel to observations on the scaling of mammalian basal metabolic rate 251 (BMR) that does not follow a single exponent across the whole BM range (Clarke 252 et al., 2010; Isaac and Carbone, 2010; Kolokotrones et al., 2010; Müller et al., 253 2012), we expect that DMI also does not follow one single exponent across the 254 whole BM range. In fact, variation in the scaling of DMI across the BM range 255 could be responsible for the observation that digesta retention scales differently in 256 different digestion types, which are distributed unevenly across the BM range 257 (Clauss et al., 2007a). In accord with Packard (2012), we do not follow the 258 argument that a curvature (or 'quadratic scaling') in double-logarithmic space

259 represents a true mechanistic pattern (Kolokotrones et al., 2010), but assume that it 260 is the effect of combined simple scaling patterns in the overall dataset and can be 261 used as a test for such scaling combinations (Müller et al., 2012). A test for such a 262 curvature effect is then followed by analysis of scaling patterns within subsets of 263 the data. In particular, following the dichotomy in metabolism scaling between 264 animals with one and with more than one offspring per litter (Müller et al., 2012), 265 we expect that food intake has a lower scaling exponent in herbivores that 266 represent the majority of the lower BM range (especially caecum fermenters), and 267 a higher scaling exponent in herbivores that represent the majority of the higher BM range (i.e. non-coprophageous hindgut fermenters, nonruminant foregut 268 269 fermenters and ruminants), resulting in a typical curvature shape in double-270 logarithmic space, or when plotting body mass-residuals of DMI against body 271 mass. An important question when assessing the individual scaling relationships of 272 intake in different groups will be whether the scaling exponent is similar to that of 273 metabolism, or whether it is actually higher (Hackmann and Spain, 2010). 274 Differences in the scaling of wet matter gut contents (WMC) and dry matter gut (v) 275 contents (DMC) with BM reflect the finding of Justice and Smith (1992) that 276 larger animals have a higher digesta moisture content; when using the WMC and 277 DMC allometries to calculate the allometry of the dry matter concentration of the 278 GIT contents, the result should be close to that of Justice and Smith (1992) where 279 DM concentration in the contents of the fermentation chamber scales to 22 BM⁻ 0.08 280 281 Species with a relatively higher food intake have a relatively shorter retention time (vi) 282 (Clauss et al., 2007a; Müller et al., 2011). Based on the above equation (6), this 283 could in theory lead to relatively lower calculated dry matter gut contents. On the 284 other hand, higher food intake itself should lead to higher calculated gut contents 285 based on the same equation (6). We aim to characterise these interdependencies in 286 our analysis. A longer relative retention time should be related to a higher relative 287 digestibility (Udén et al., 1982; Clauss et al., 2007b). 288 Because one of the largest existing datasets on the digestive physiology of large herbivores

200 Decause one of the furgest existing datasets on the digestive physiology of furge herorvores

289 (Foose, 1982) is basically incompatible with other published data (Clauss et al., 2007a; Clauss

et al., 2010b), we performed the same analyses for both the Foose (1982) dataset and an

291 independent collation of literature data. With the limitation that the Foose (1982) dataset

comprises only comparatively large species, this allows a corroboration of results from onedataset with the other.

294

295 Methods

296 The datasets used in this study are

- a) Foose (1982), comprising 26 species of large (>100 kg) herbivores from the
- 298 299

300

Artiodactyla, Perissodactyla and Proboscidea, which received two standardized diets (grass and lucerne hay); in this dataset, the unit of reference is organic matter rather than dry matter; and

301 b) the data collation of Clauss et al. (2007a) that was supplemented with additional 302 sources, and pruned according to the following selection criteria. Only sources were 303 used which gave body mass (BM), dry matter intake (DMI), particle mean retention time in the whole gastrointestinal tract (MRT_{particle}GIT,) and the apparent digestibility 304 305 of dry matter (aD DM) from the same experiment (in some cases, this information had 306 to be collected from several publications reporting on the same experiment). From 307 these data, dry matter content (DMC) of the GIT was calculated as outlined in the 308 Indtroduction. For the sake of a large data collection, we did not select publications 309 according to the passage markers used. Therefore, any particle marker was accepted; 310 however, only studies in which small particles (generally ≤ 2 mm; as opposed to whole forages) were labelled were included. Similarly, we did not select publications 311 312 according to the type of food offered; therefore, the data collection comprises animals 313 fed on a complete, pelleted diet, animals on mixtures of roughages and concentrates, 314 as well as animals on roughage-only diets. Ideally, a data collection with a 315 standardized diet (such as in Foose, 1982; Steuer et al., 2011; 2012) or with diets 316 representing the natural diets of the species would be preferable, but are not available 317 at a larger scale and over a wide body mass range. With regards to some domestic 318 species (rabbits, horse, goats, sheep, cattle), no attempt was made to collate all 319 published data. For lagomorphs and rodents, only data from trials in which 320 coprophagy was not prevented by the study design were used. For each species, one 321 average value for BM, DMI, MRT_{particle}GIT, aD DM and DMC was calculated using 322 all available data from the cited publications. The data for the resulting 77 mammalian 323 species and its sources are summarized in the Appendix.

324 c) To compare the relationship of wet matter contents (WMC) of the GIT with the dry
 325 matter contents, the data collection on WMC collated in Clauss et al. (2007a; n=72
 326 species) was used.

327 Species were allocated, according to Stevens and Hume (1995), to one of the following 328 categories: caecum fermenters, colon fermenters, non-ruminant foregut fermenters, and 329 ruminants (Appendix). Additionally, species were allocated to litter size classes (one or more 330 than one offspring per litter) and body size classes (<10 kg, >10 kg). In order to account for 331 ancestry-biased correlations in the datasets (i.e., significant associations in interspecific 332 datasets might be due to the fact that phylogenetically related species exhibit similar eco-333 physiological traits; Harvey and Pagel, 1991), the data were controlled for phylogenetic 334 influences using the "Phylogenetic Generalized Least-Squares" method (PGLS; Martins and 335 Hansen, 1997; Rohlf, 2001). This procedure estimates a covariance matrix of the species due 336 to their ancestral roots and includes these interrelationships as error term in a generalized least 337 squares algorithm to determine the model parameters. The phylogenetic trees for the two 338 datasets were derived by pruning the mammal supertree from Bininda-Emonds et al. (2007,

- 339 2008) of those species not represented in the datasets.
- 340

341 Allometric regressions were performed as linear regressions on log-transformed data, with

342 digestive parameters (DMI, MRT_{particle}GIT, aD DM, DMC) as the respective dependent

343 variables, and with log-transformed BM as the independent variable. Tests for a 'curvature' in

344 double-logarithmic space were made using quadratic regressions. Statistical analyses were

345 performed without (ordinary least squares regressions: OLS) and with accounting for

phylogeny (PGLS), in PASW 18.0 (SPSS Inc., Chicago, IL) and in *R* 2.15.0 (Team, 2011)

347 with the caper package. The significance level was set to α =0.05.

348

349 **Results**

350 Foose (1982) dataset

351 Variables scaled as: Food intake (as organic matter intake, OMI) with BM^{0.76}, apparent

digestibility (of organic matter) with $BM^{0.02}$, organic matter gut contents (OMC) with $BM^{0.82}$,

and digesta retention with $BM^{0.07}$, which matches the predicted relationships of 0.07 ~ [0.82 –

354 0.76 + 0.02] (Table 1). Notably, the 95% confidence interval for the scaling of OMC did not

include 1.00 (Table 1). Results were similar in PGLS. Similar patterns were evident within

the hindgut fermenters and the ruminants only (Table 1). Both digesta retention and gut fill

357 had a scaling exponent lower than expected.

358

359	Within the body mass residuals, intake was negatively correlated with digesta retention in the
360	whole dataset in OLS but not in PGLS (Table 2), indicating that certain taxonomic groups
361	(notably the ruminants) did not follow this pattern (Fig. 1a). Intake was positively correlated
362	with gut fill in both OLS and PGLS (Table 2, Fig. 1b). There was a positive correlation
363	between digesta retention and digestibility in OLS but not in PGLS (Table 2), because these
364	two residuals showed a clear dichotomy between ruminants and colon fermenters (Fig. 1c).
365	Digesta retention was not correlated with gut fill in OLS but it was in PGLS (Table 2),
366	indicating that the relationship only existed within more closely related groups (Fig. 1d).
367	
368	Our dataset
2 ()	\mathbf{M} is a property of \mathbf{M} of \mathbf{M} is a property of \mathbf{M} of \mathbf{M}

Variables scaled as: Food intake (as dry matter intake, DMI) with BM^{0.76} (Fig. 2a), apparent 369

digestibility (of dry matter, aD DM) with BM^{-0.03} (Fig.2c), dry matter gut contents (DMC) 370

with BM^{0.93} (Fig. 2e), and digesta retention with BM^{0.16} (Fig. 2g) (all in OLS), which matches 371

372 the predicted relationships of $0.16 \sim [0.93 - 0.76 + (-0.03)]$. Notably, the 95% confidence

373 interval for the scaling of DMC did not include 1.00 (Table 3). Results for PGLS were

similar; notably, the scaling exponent for MRT was lower than in OLS at $BM^{0.12}$. 374

375

376 When testing for a curvature effect in the overall dataset in OLS, such a curvature (visible 377 when plotting body mass residuals against body mass) was evident for dry matter intake (Fig. 378 2b) and for digesta retention (Fig. 2h), but not for digestibility (Fig. 2d) or gut contents (Fig. 379 2f). The full term describing the curvature was significant, with 95% confidence intervals that 380 did not include zero for the quadratic term, for intake and digesta retention only (Table 4); the 381 exponent of the quadratic term had a similar magnitude but a different sign for these two 382 measurements (intake: 0.053, retention: -0.057). In PGLS, these quadratic terms were not 383 significant (Table 4), indicating that the effect of the curvature was caused by the

384 phylogenetic structure of the dataset.

385

386 Splitting the overall dataset led to different scaling exponents for the various subsets (Table 387 5). The scaling of food intake had a lower exponent among species that produce more than 388 one offspring per litter or that are below a 10 kg threshold (95%CI 0.57-0.70 in OLS and 389 0.64-0.81 in PGLS), as opposed to species with one offspring per litter or above the 10 kg

390 threshold (95%CI 0.82-0.96 in OLS and 0.74-0.95 in PGLS). The scaling of dry gut contents

391 was similar among these groups (0.91-0.94 in OLS and 0.84-0.96 in PGLS; never including 392 1.00 in the 95%CI in OLS but for the larger animals in PGLS). In OLS, there was a difference 393 in the scaling of digesta retention between these groups, with a significant scaling exponent in 394 animals with more than one offspring per litter or below the 10 kg threshold, in contrast to no 395 significant scaling (95% confidence interval for the exponent included zero) in animals above 396 the 10 kg threshold (Table 5). In PGLS, the exponent of the MRT scaling in the animals with 397 more than one offspring per litter/below the 10 kg threshold was distinctively lower than in 398 OLS, with 95%CI approaching or including zero (Table 5) and thus making the scaling in this group similar to the scaling in the group of animals with one offspring per litter/more than 10 399 400 kg. Notably, the 95%CI of the scaling exponent of intake and dry gut contents overlapped in 401 animals with a single offspring/animals >10 kg in OLS, and for all herbivores in PGLS.

402

403 Splitting the species into the four digestion types resulted in similar patterns, with a lower 404 scaling exponent for intake, and a higher one for digesta retention, in the (small) caecum 405 fermenters as compared to the other groups in OLS (Table 6). In PGLS, these differences 406 were not evident; in particular, the scaling exponent for MRT in caecum fermenters was much 407 lower in PGLS at 0.14 than in OLS at 0.27, suggesting the high scaling previously reported in 408 caecum fermenters of 0.25 (Clauss et al., 2007a) was due to an uneven distribution of 409 phylogenetically distant species within this data subset (Table 6). A scaling exponent of 1.00 410 for gut fill was only included in the confidence interval for ruminants in OLS, and for colon 411 fermenters and ruminants in PGLS. The 95% CI of the scaling exponent overlapped for intake 412 and gut fill in all but the caecum fermenters in OLS, and in all four groups in PGLS (Table 6). 413

415

414 Within the body mass residuals of the whole dataset (Table 7), intake was negatively

415 correlated with digesta retention (Fig. 3a) and positively with gut fill (Fig. 3b) in both OLS

and PGLS. Although digestibility was negatively related to intake and positively related to

417 digesta retention (Fig. 3c) in OLS, these relationships were not significant in PGLS (Table 7),

418 i.e. they do not occur within more closely related taxa. By contrast, digestibility was

419 negatively related to gut fill in PGLS, indicating that among more closely related species,

420 those with a higher gut fill had lower digestibilities. Gut fill was positively related to digesta

421 retention in both OLS and PGLS (Table 7, Fig. 3d).

422

423 Wet gut contents

424 When testing for a scaling curvature in the dataset on wet gut contents, the quadratic term was

425 not significant (OLS p=0.706; PGLS p=0.463). Wet gut contents scaled to 0.108 (0.095-

426 0.123) BM^{1.062(1.029-1.094)} (R²=0.984, p<0.001) in OLS and 0.109 (0.036-0.332) BM^{1.086(1.004-1.004)}

427 $^{1.167)}$ (R²=0.907, p<0.001) in PGLS. Thus, when comparing the scaling of wet gut contents 428 and of dry matter gut contents, the difference between the two increases with body mass (Fig.

429 4); from the comparison of the two scaling exponents (OLS 0.93 and 1.06; PGLS 0.92 and

- 430 1.09), the dry matter concentration of gut contents should theoretically scale to BM^{-0.13} or
- 431 BM^{-0.17}.
- 432

433 **Discussion**

434 The aim of this study was to test whether empirical data allow evoking differences in the 435 allometric scaling between physiological characteristics, to explain species diversification and 436 niche differentiation along a body size gradient in herbivores. Two effects have previously 437 been associated with body size-driven diversification in mammalian herbivores, namely a 438 higher scaling of food intake as compared to the scaling of metabolism (Hackmann and Spain, 439 2010), and a distinct difference in the scaling of food intake and gut fill, leading to a positive 440 scaling of digesta retention time (Demment and Van Soest, 1985). If we assume that the 441 results of controlled feeding studies are indeed representative of the digestive physiology of 442 the animals investigated, then the results indicate that from these two effects, the former 443 represents the more promising approach, in particular for larger herbivores. 444

445 We suggest several reasons for the fact that the concept of an increasing difference between 446 intake and gut fill, and hence increasing digesta retention time with body size, does not apply 447 to herbivores to the extent previously suggested. Indeed, the concept has been questioned 448 repeatedly based on the empirical finding that the expected scaling of digesta retention with 449 body mass cannot be demonstrated (see Introduction). Similarly, the results of our study 450 demonstrate that any scaling of digesta retention, though principally existent, is below the 451 quarter-power scaling often associated with biological times (Illius and Gordon, 1992; Brown 452 et al., 2012). Because of the mathematical relationships between intake, digestibility, retention 453 and gut fill, this translates into the following deviations from the original concept: gut fill 454 does not scale linearly with body mass, but to a slightly lower exponent (the 95%CI often did 455 not include 1.0); on the other hand, intake scales to an exponent higher than 0.75 in large 456 herbivores. Therefore, the difference between the two scaling exponents is lower than 457 assumed when interpreting the Jarman-Bell principle in terms of digestive physiology, with 458 confidence intervals overlapping in many datasets. Additionally, there is a slight negative 459 scaling of digestibility with body mass in these experiments, which also reduces the scaling

460 exponent of retention time (see Introduction). The ultimate reason why digesta retention

shows a low scaling with body mass in herbivores might stem from the fact that from a

462 certain body size upwards, a further increase in retention time will not, on average, convey

463 additional advantages, principally because the additional gain from exposing digesta to the

464 digestive process for a longer time is lower than the gain from additional food intake

465 (Hummel et al., 2006).

466

Several arguments can be put forward to support the validity of our findings. The below-467 468 linear scaling of gut contents was present in both of the datasets investigated (Tables 1 and 3). 469 The assumption that gut contents (as a proxy for gut capacity) scale linearly with body mass 470 was originally derived from datasets that related wet gut content mass to body mass (Parra, 471 1978; Demment and Van Soest, 1985; Justice and Smith, 1992; Clauss et al., 2007a; Illius and 472 Gordon 1992 is the only publication that uses dry matter gut contents and also arrives at linear 473 scaling, but the dataset comprised only 16 species). If one assumes that this finding translates 474 directly into a similar scaling for that unit of gut fill related to food intake (i.e. dry matter), 475 then one makes the silent assumption that the proportion of moisture in gut contents is similar 476 across all body sizes (as done by Demment, 1983). However, Justice and Smith (1992) 477 reported that the DM concentration of the contents of the fermentation chamber scales to BM⁻ $^{0.08}$ – in other words, the contents of this part of the gut becomes more watery in larger 478 479 animals. Their finding is corroborated by our comparison of the scaling of total wet and dry matter gut contents, which yields a similar negative scaling at BM^{-0.13} to BM^{-0.17}. Similarly, 480 Müller et al. (2011) found, in a comparative analysis of mean retention times of particle and 481 482 solute (fluid) markers, that the difference between these two digesta phases tends to increase 483 with body mass – not necessarily because particles are retained longer in larger animals, but 484 also because fluids are passed through their digestive system at a higher rate. One non-485 exclusive ultimate reason for this might be the fact that in the larger guts of larger animals, 486 diffusion distances from within the digesta to the sites of absorption in the gut wall are 487 increased (Clauss and Hummel, 2005), a problem that could be solved by a more fluid digesta 488 (Lentle et al., 2006). Thus, we suggest that the assumption that that part of gut fill relevant to 489 food intake and digestibility (i.e. the dry matter gut fill) scales linearly with body mass cannot 490 be defended until more data become available. 491

The other important assumption that is challenged by our findings is that food intake scales
uniformly to BM^{0.75}. Unfortunately, a direct inclusion of data on basal metabolic rate (BMR)

494 in our study is not feasible; although BMR has been measured in a large variety of 495 mammalian species (McNab, 2008), the large animals that form a majority of our dataset are 496 not included in the available BMR data pool. Therefore, we need to refer to comparisons of 497 our findings to those from BMR studies, inherently with a distinct difference in the species 498 sets investigated. In the debate on the scaling of basal metabolic rate, it has been found 499 repeatedly that across the whole mammalian body mass range, a single allometric exponent 500 does not represent the data very precisely, but that either a curvature in log-log space, or a 501 combination of different scaling relationships of subgroups provides a better fit (Hayssen and 502 Lacy, 1985; Clarke et al., 2010; Isaac and Carbone, 2010; Kolokotrones et al., 2010; Müller et 503 al., 2012). The magnitude of an overall mammalian scaling exponent will therefore depend 504 either on the body mass range included in the analysis (Lovegrove, 2000; Dodds et al., 2001; 505 Glazier, 2005; White and Seymour, 2005), the taxonomic composition of the sample 506 (Hayssen and Lacy, 1985; Sieg et al., 2009; White et al., 2009; Capellini et al., 2010), or on 507 the inclusion of animals according to certain biological characteristics (McNab, 2008; Müller 508 et al., 2012). Although our species sample of mammalian herbivores is lower than the datasets 509 used to investigate the scaling of metabolism, a similar 'curvature effect' is evident in the 510 intake and retention data (Fig. 2ab and 2gh). Our findings thus provide further support to 511 these previous studies that scaling exponents are not necessarily a universal characteristic for 512 the whole mammalian clade, but may differ either between different groups, or vary across 513 the body mass range. In particular, the results indicate that the group of mammals that can be 514 classified as either small, giving birth to multiple young at a time, or being caecum fermenters 515 has a generally lower scaling of food intake than animals classified as larger, giving birth to a 516 single offspring, or being colon fermenters, nonruminant foregut fermenters or ruminants. 517 Which of these classifications is biologically relevant for the difference in scaling, or whether 518 they are all proxies for a yet undefined functional subdivision of mammals, remains to be 519 investigated. Because a similar 'curvature effect' is evident in the scaling of metabolic rate, 520 where not only herbivores but also other feeding types were included in the dataset 521 (Kolokotrones et al., 2010; Müller et al., 2012), we suggest that digestive strategy is unlikely 522 to be the relevant proxy. Regardless, our study suggests that the use of one single exponent to 523 describe relative metabolic rates or relative food intakes may be convenient, but should not be 524 the basis for biological models.

525

Within one notable subset of large herbivores, the ruminants, the assumption that food intake
 scales to BM^{0.75} has previously been refuted based either on conceptual approaches or

collections of empirical data: Mertens (1994) and Südekum (2002) explain that for low-528 529 quality forages, relating food intake in domestic ruminants to body mass may be more 530 appropriate than using metabolic body mass as the basis. Hackman and Spain (2010) collated 531 data on the voluntary food intake (VFI) in domestic and wild ruminant species in a dataset 532 more comprehensive than ours (that was limited to studies that also measured digestibility and 533 retention), and concluded that VFI scaled to BM^{0.9}, which was similar to that described in 534 previous studies on domestic ruminants (Graham, 1972; Minson, 1990; Reid et al., 1990). The 535 authors concluded that the intermediate scaling between metabolism (0.75) and gut fill (1.0)536 indicated that both metabolic and physical constraints determine food intake; they also 537 conclude that the putative difference between the scaling of metabolism (0.75) and the scaling 538 of intake (0.9) allows larger animals to ingest lower-quality food and still meet their energetic 539 requirements. Note that this interpretation of scaling differences does not draw on an alleged 540 scaling of digestion efficiency and retention times.

541

542 The assumption that intake might scale to a higher exponent than basal metabolic rate is 543 supported when comparing the intake scaling exponent of large herbivores/herbivores with 544 one offspring of this study (0.81-0.89, Table 5) with that of the BMR in mammals with one 545 offspring (0.76; Müller et al. 2012). In contrast, for the small herbivores/herbivores with more 546 than one offspring, such a difference is not evident at 0.62-0.72 for the intake scaling 547 exponent in this study and 0.69 for that of BMR (Müller et al. 2012). To explain body size 548 effects in this group, factors such as variation in feeding selectivity and caecotroph use via 549 coprophagy must be further investigated (Justice and Smith, 1992; Franz et al., 2011b). In 550 particular, it remains unknown whether the proportion of intake achieved via coprophagy 551 varies systematically with body mass. Preliminary results indicate that this proportion 552 increases with increasing dietary fibre content (Takahashi and Sakaguchi, 1998). If we 553 assume that compared to smaller species, larger caecum fermenters are adapted to lower-554 quality diets (i.e. diets higher in fibre), and should therefore be generally adapted to a higher 555 proportion of intake via coprophagy, this might help explain the lower scaling of intake via 556 food (as measured in the digestion experiments collated in this study) in this group. Further 557 studies that assess the contribution of coprophagy over a wide range of body sizes are 558 required to explore this possibility.

559

560 Apart from the logical issue with drawing conclusions on the scaling of digestibility from the 561 scaling of digesta retention explained in the Introduction, our data collection yields some further insights concerning correlates and effects of digesta retention. As reported previously in inter- and intraspecific analyses (Clauss et al., 2007a; Clauss et al., 2007b; Müller et al., 2011; Steuer et al., 2011), species with a higher relative food intake have comparatively shorter retention times (Fig. 3a). Langer and Snipes (1991) suggested that interspecifically, animals achieve longer digesta retention by higher gut capacities, a concept supported by our findings (Fig. 3b). Additionally, capacity for a higher gut fill will allow a higher food intake (Fig. 3d).

569

570 Although increasing retention times may be associated with higher general digestive 571 efficiencies in intraspecific studies (Clauss et al., 2007b), this effect is most evident in 572 interspecific comparisons in relation to the digestibility of the fibre component of the food 573 (Udén and Van Soest, 1982; Clauss et al., 2009). When comparing the overall diet 574 digestibility, as in this study, where we collated data on the dry matter or organic matter 575 digestibility but not the fibre digestibility, no significant effect of digesta retention on 576 digestibility is evident when accounting for the phylogenetic structure of the data (Table 3 and 577 7). In other words, although larger clades differ systematically in retention and digestibility, 578 such as the colon fermenters and ruminants in Fig. 1c, the effect is not evident among more 579 closely related species – in the case of the Foose (1982) dataset even though consistent diets 580 were used. One potential reason is that digesta retention is not the only determinant of 581 digestibility, but acts together with digesta particle size reduction, either compensating for 582 poor chewing efficiency (as for example in hippopotamuses) or reinforcing a high chewing 583 efficiency (as in the ruminants) (Clauss et al., 2009). Another potential reason is that overall 584 dry matter digestibility does not necessarily reflect fibre digestibility. A small-scale data 585 collection in Clauss and Hummel (2005) suggested that fibre digestibility may vary more 586 across a large herbivore body mass range than does overall diet digestibility, which remained 587 rather constant (as in this study). The authors summarized concepts that interpret these data as 588 indication for a better utilization of non-fibrous food components in smaller animals, 589 including the segregation of bacterial protein during caecotroph formation by the colonic 590 separation mechanism (Björnhag, 1987; Franz et al., 2011b). Understanding how bacterial 591 nitrogen utilization and endogenous/metabolic nitrogen losses vary with body mass and 592 between the different herbivore digestive strategies remains one of the major challenges in 593 studying herbivore digestive physiology (Schwarm et al., 2009b).

595 Herbivores will maximize energy intake if they maximize feeding selectivity, food intake and 596 digesta retention (by increasing gut fill), particle size reduction (by more elaborate cranio-597 dental designs and by sorting mechanisms in ruminants that increase chewing efficiency), and 598 minimize endogenous/metabolic faecal losses. The major dilemma in this array of factors is 599 that increasing food intake is usually associated with a lower feeding selectivity, shorter 600 digesta retention and reduced particle size reduction (Clauss et al., 2010a). It is commonly 601 assumed that the different herbivore species or groups represent different, equally successful 'digestive strategies' to solve this dilemma, varying in the emphasis of the individual 602 603 interrelated factors (Hume, 2005). Alternatively, one can hypothesize that the sequence of 604 radiation events in herbivores indicates different levels of optimization of the overall interplay 605 of these factors (Fritz et al., 2009). For example, bovid ruminants, as the most recently 606 radiated large herbivore group, potentially represent the hitherto most sophisticated design 607 that combines high food intakes, long retention times and high chewing efficiency with a very 608 efficient use of the symbiotic microflora (Clauss et al., 2010a; Müller et al., 2011). How 609 species that use similar niches with different digestive strategies can persist then requires 610 different explanations. Such a view shifts the focus in tracing niche differentiation and species 611 diversification from simple allometric considerations to more complex aspects of variation in 612 organismal design.

613

614 Acknowledgements

615 This work was supported by DFG grants CL 182/3-1, 5-1 and 6-1. This is contribution no.

616 119 of the DFG Research Group 533 Biology of the Sauropod Dinosaurs: The Evolution of

- 617 Gigantism.
- 618

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У	stats	a	95%CI	b	95%CI	\mathbf{r}^2	р
			all spec	ies (n=26)			
OMI		0.064	0.027-0.153	0.759	0.621-0.897	0.843	<0.001
aDOM	OLS	53.6	40.6-70.6	0.015	-0.029-0.059	0.020	0.494
OMC	OLS	0.074	0.038-0.147	0.819	0.710-0.927	0.910	< 0.001
MRT		38.2	22.2-65.6	0.065	-0.020-0.151	0.094	0.128
OMI		0.052	0.024-0.114	0.792	0.690-0.895	0.905	<0.001
aDOM	DOL 0	45.7	34.7-60.2	0.033	-0.004-0.069	0.115	0.062
OMC	PGLS	0.063	0.025-0.161	0.842	0.719-0.964	0.883	<0.001
MRT		37.3	21.6-64.5	0.062	-0.010-0.134	0.107	0.075
			Hindgut fer	menter (n=	11)		
OMI		0.097	0.034-0.281	0.726	0.564-0.888	0.920	<0.001
aDOM	OLS	46.5	36.1-59.7	0.024	-0.015-0.062	0.180	0.194
OMC		0.085	0.037-0.196	0.809	0.681-0.937	0.958	<0.001
MRT		27.1	16.9-43.6	0.092	0.019-0.164	0.477	0.019
OMI		0.083	0.029-0.242	0.748	0.597-0.899	0.913	<0.001
aDOM	PGLS	37.1	26.9-51.3	0.054	0.008-0.099	0.375	0.029
OMC	PGLS	0.074	0.026-0.208	0.824	0.679-0.970	0.932	<0.001
MRT		25.2	14.9-42.8	0.098	0.023-0.172	0.425	0.017
			Rumina	int (n=13)			
OMI		0.104	0.029-0.372	0.664	0.452-0.875	0.813	<0.001
aDOM	OLS	52.1	33.8-80.2	0.031	-0.040-0.103	0.079	0.353
OMC	ULS	0.126	0.028-0.581	0.728	0.476-0.980	0.786	<0.001
MRT		39.3	17.1-89.7	0.079	-0.058-0.216	0.128	0.231
OMI		0.099	0.034-0.283	0.665	0.504-0.825	0.857	<0.001
aDOM	DCLO	67.2	43.8-103.0	-0.010	-0.075-0.055	0.008	0.911
OMC	PGLS	0.165	0.037-0.737	0.687	0.460-0.915	0.761	<0.001
MRT		61.0	23.7-156.9	0.016	-0.128-0.160	0.004	0.954

Table 1. Scaling relationships of parameters of digestive physiology with body mass (BM) in the Foose (1982) dataset using OLS and PGLS statistics according to $y = a BM^b$.

organic matter intake (OMI in kg d⁻¹); apparent digestibility of organic matter (aDOM in %);
organic matter gut contents (OMC in kg); particle mean retention time (MRT in h)

858

859 Table 2. Correlations between body mass residuals of parameters of digestive physiology in

all species of the Foose (1982) dataset analysed by OLS and PGLS statistics (differences

861 indicated by grey shading).

	stats	Residual aDOM	Residual OMC	Residual MRT
Residual OMI	OLS	R=-0.137, p=0.505	R=0.841, p<0.001	R=-0.577, p=0.002
	PGLS	R=-0.300, p=0.114	R=0.828, p<0.001	R=0.033, p=0.975
Residual aDOM	OLS		R=0.173, p=0.398	R=0.654, p<0.001
	PGLS		R=0.297, p=0.120	R=0.293, p=0.126
Residual OMC	OLS			R=-0.055, p=0.790
	PGLS			R=0.575, p<0.001

organic matter intake (OMI in kg d⁻¹); apparent digestibility of organic matter (aDOM in %);
organic matter gut contents (OMC in kg); particle mean retention time (MRT in h)

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Table 3. Scaling relationships of parameters of digestive physiology with body mass (BM) in the dataset of this study (n=77) using OLS and PGLS statistics according to $y = a BM^b$.

		10 00000) (11 11) 451				0
У	stats	a	95%CI	b	95%CI	r ²	р
DMI		0.046	0.041-0.052	0.763	0.732-0.794	0.970	< 0.001
aDDM	OI C	65.5	61.9-69.2	-0.025	-0.0390.011	0.147	0.001
DMC	OLS	0.029	0.025-0.032	0.931	0.902-0.960	0.982	< 0.001
MRT		22.3	19.5-25.5	0.158	0.124-0.191	0.540	< 0.001
DMI		0.039	0.017-0.089	0.773	0.720-0.825	0.916	< 0.001
aDDM	DCLC	67.8	42.3-108.6	-0.052	-0.0820.021	0.130	< 0.001
DMC	PGLS	0.028	0.010-0.075	0.915	0.851-0.979	0.913	< 0.001
MRT		26.1	11.639-58.4	0.120	0.068-0.172	0.214	< 0.001

dry matter intake (DMI in kg d⁻¹); apparent digestibility of dry matter (aDDM in %); dry
 matter gut contents (DMC in kg); particle mean retention time (MRT in h)

870 871

872 Table 4. Evidence for quadratic scaling in parameters of digestive physiology with body mass

873 (BM) in the dataset of this study (n=77) (OLS). Linear regression models according to

у	stats	a	95%CI	b1	95%CI	b2	95%CI	R2 (adj.)	р
DMI		0.042	0.038-0.047	0.659	0.616-0.702	0.053	0.035-0.070	0.980 (0.979)	<0.00
aDDM	OI S	66.7	63.1-70.5	-0.004	-0.027-0.018	-0.010	-0.0190.001	0.202 (0.181)	<0.00
DMC	OLS	0.028	0.025-0.032	0.930	0.881-0.979	0.001	-0.019-0.021	0.982 (0.981)	<0.00
MRT		24.7	22.0-27.7	0.270	0.223-0.316	-0.057	-0.0760.038	0.692 (0.983)	<0.00
DMI		0.039	0.018-0.088	0.706	0.617-0.795	0.026	-0.002-0.054	0.920 (0.918)	<0.00
aDDM	PGLS	67.7	42.1-108.9	-0.056	-0.1080.003	0.001	-0.015-0.018	0.130 (0.106)	0.0
DMC	POLS	0.028	0.010-0.074	0.851	0.743-0.959	0.025	-0.009-0.060	0.915 (0.913)	<0.00
MRT		26.1	11.6-58.7	0.121	0.032-0.211	-0.001	-0.029-0.028	0.214 (0.193)	<0.00

874 $\log y = a + b1(\log BM) + b2((\log BM)^2)$ or $y = a BM^{(b1+b2(\log BM))}$

875 dry matter intake (DMI in kg d⁻¹); apparent digestibility of dry matter (aDDM in %); dry

876 matter gut contents (DMC in kg); particle mean retention time (MRT in h)

Table 5. Scaling relationships of parameters of digestive physiology with body mass (BM) in

У	stats	a	95%CI	b	95%CI	r^2	р
			Litter size	> 1 (n=24)		
DMI		0.050	0.046-0.055	0.663	0.626-0.701	0.984	<0.00
aDDM	OI C	62.230	57.4-67.3	-0.023	-0.054-0.008	0.095	0.142
DMC	OLS	0.027	0.022-0.034	0.905	0.825-0.986	0.961	<0.00
MRT		19.275	16.0-23.2	0.233	0.160-0.307	0.662	<0.00
DMI		0.047	0.032-0.067	0.724	0.647-0.802	0.938	<0.00
aDDM	DOLG	64.3	50.7-81.6	-0.015	-0.066-0.035	0.015	0.712
DMC	PGLS	0.028	0.014-0.058	0.884	0.733-1.036	0.856	< 0.00
MRT		21.8	12.6-37.9	0.153	0.036-0.270	0.230	0.00
			Litter size	= 1 (n=53)		
DMI		0.029	0.024-0.036	0.864	0.818-0.910	0.966	<0.00
aDDM	01.0	72.611	65.9-80.0	-0.046	-0.0670.024	0.260	<0.00
DMC	OLS	0.028	0.023-0.035	0.935	0.891-0.978	0.973	<0.00
MRT		36.475	30.3-44.1	0.051	0.009-0.093	0.105	0.01
DMI		0.030	0.013-0.074	0.814	0.747-0.881	0.917	<0.00
aDDM DMC	DCLG	67.0	45.0-99.8	-0.040	-0.0710.010	0.117	0.00
	PGLS	0.026	0.010-0.070	0.916	0.841-0.990	0.919	<0.00
MRT		31.5	14.1-70.2	0.084	0.023-0.145	0.125	0.002
			<10 kg	(n=36)			
DMI		0.045	0.040-0.050	0.622	0.568-0.677	0.941	<0.00
aDDM	OLS	67.453	64.3-70.8	0.020	-0.005-0.045	0.070	0.11
DMC	ULS	0.028	0.024-0.032	0.909	0.834-0.984	0.947	<0.00
MRT		22.751	19.8-26.2	0.297	0.223-0.371	0.661	<0.00
DMI		0.038	0.021-0.069	0.723	0.640-0.806	0.896	<0.00
aDDM	PGLS	67.3	49.7-91.2	0.007	-0.036-0.050	0.003	0.89
DMC	TOLS	0.029	0.011-0.074	0.840	0.707-0.973	0.819	<0.00
MRT		27.4	11.4-65.9	0.119	-0.005-0.243	0.095	0.04
			> 10 kg	g (n=41)			
DMI		0.026	0.018-0.037	0.885	0.815-0.955	0.944	< 0.00
aDDM	OLS	64.863	53.6-78.5	-0.029	-0.066-0.009	0.058	0.12
DMC		0.030	0.021-0.043	0.924	0.854-0.993	0.948	< 0.00
MRT		41.687	29.9-58.2	0.026	-0.038-0.091	0.017	0.41
DMI		0.029	0.010-0.081	0.844	0.743-0.945	0.873	<0.00
aDDM	PGLS	54.8	34.6-86.7	-0.003	-0.049-0.042	0.001	0.97
DMC	I ULS	0.023	0.007-0.070	0.959	0.848-1.070	0.880	< 0.00
MRT		26.3	12.1-57.5	0.114	0.037-0.190	0.176	0.00

the dataset of this study in subsets according to litter size or a 10 kg body mass threshold using OLS and PGLS statistics according to $y = a BM^b$.

dry matter intake (DMI in kg d⁻¹); apparent digestibility of dry matter (aDDM in %); dry
matter gut contents (DMC in kg); particle mean retention time (MRT in h)

885Table 6. Scaling relationships of parameters of digestive physiology with body mass (BM) in886the dataset of this study in subsets according to digestion types using OLS and PGLS statistics887according to $y = a BM^b$.

у	stats	a	95%CI	b	95%CI	r ²	р
			Caecum ferr	menter (n=	=25)		
DMI		0.041	0.036-0.048	0.589	0.524-0.654	0.938	<0.001
aDDM	OLS	65.3	61.4-69.5	0.001	-0.029-0.031	0.000	0.950
DMC	OLS	0.025	0.021-0.031	0.860	0.767-0.954	0.940	<0.001
MRT		21.7	17.8-26.5	0.271	0.177-0.365	0.606	<0.001
DMI		0.033	0.017-0.066	0.688	0.594-0.782	0.900	<0.001
aDDM	PGLS	67.2	50.7-89.2	-0.003	-0.042-0.036	0.001	0.977
DMC	POLS	0.025	0.009-0.070	0.830	0.689-0.972	0.852	<0.001
MRT		27.4	11.5-65.1	0.140	0.021-0.258	0.187	0.013
			Colon ferm	nenter (n=1	15)		
OMI		0.024	0.018-0.033	0.911	0.856-0.967	0.990	<0.001
DDM	OLS	78.7	61.9-100.0	-0.066	-0.1090.023	0.459	0.006
OMC	OLS	0.029	0.021-0.040	0.919	0.861-0.977	0.989	<0.001
MRT		47.1	27.2-81.5	-0.021	-0.120-0.077	0.017	0.647
DMI		0.031	0.016-0.058	0.852	0.757-0.947	0.960	<0.001
aDDM DMC	DCLS	75.7	49.3-116.1	-0.070	-0.1340.006	0.259	0.032
	PGLS	0.027	0.015-0.049	0.948	0.858-1.037	0.971	<0.001
MRT		34.4	14.3-82.6	0.061	-0.071-0.192	0.060	0.461
		No	onruminant foreg	gut fermen	ter (n=15)		
DMI		0.036	0.026-0.051	0.755	0.655-0.855	0.953	<0.001
aDDM	OLS	69.8	56.0-87.3	-0.038	-0.105-0.030	0.102	0.246
DMC	OLS	0.030	0.023-0.038	0.881	0.806-0.957	0.980	<0.001
MRT		30.3	21.4-43.0	0.109	0.003-0.215	0.274	0.045
DMI		0.031	0.009-0.113	0.812	0.659-0.965	0.893	<0.001
aDDM	PGLS	75.1	46.7-120.9	-0.064	-0.1200.008	0.276	0.025
DMC	TOLS	0.033	0.009-0.114	0.844	0.697-0.992	0.907	<0.001
MRT		40.2	11.1-145.1	0.006	-0.145-0.158	0.001	0.993
			Rumina	nt (n=22)			
DMI		0.054	0.035-0.083	0.752	0.659-0.845	0.934	<0.001
aDDM	OLS	66.7	51.6-86.3	-0.025	-0.080-0.030	0.044	0.347
DMC	CLO	0.041	0.024-0.067	0.897	0.788-1.005	0.937	<0.001
MRT		27.6	20.2-37.7	0.133	0.066-0.200	0.464	<0.001
DMI		0.036	0.014-0.093	0.765	0.660-0.870	0.910	<0.001
aDDM	PGLS	85.3	42.8-169.9	-0.072	-0.149-0.005	0.144	0.055
DMC		0.024	0.007-0.078	0.973	0.841-1.104	0.913	<0.001
MRT		26.3	13.2-52.5	0.177	0.100-0.254	0.502	<0.001

dry matter intake (DMI in kg d⁻¹); apparent digestibility of dry matter (aDDM in %); dry
 matter gut contents (DMC in kg); particle mean retention time (MRT in h)

- Table 7. Correlations between body mass residuals of parameters of digestive physiology in 893
- the dataset of this study (n=77) analysed by OLS and PGLS statistics (differences indicated 894 by grev shading). 895

by grey shading	.)•			
	Stats	Residual aD DM	Residual DMC	Residual MRT
Residual DMI	OLS	R=-0.254, p=0.026	R=0.444, p<0.001	R=-0.583, p<0.001
	PGLS	R=0.087, p=0.567	R=0.589, p<0.001	R=-0.270, p=0.004
Residual aD DM	OLS		R=-0.082, p=0.471	R=0.345, p=0.002
	PGLS		R=-0.200, p=0.050	R=-0.084, p=0.592
Residual DMC	OLS			R=0.450, p<0.001
	PGLS			R=0.593, p<0.001
		1		

dry matter intake (DMI in kg d^{-1}); apparent digestibility of dry matter (aDDM in %); dry 896 897

matter gut contents (DMC in kg); particle mean retention time (MRT in h)

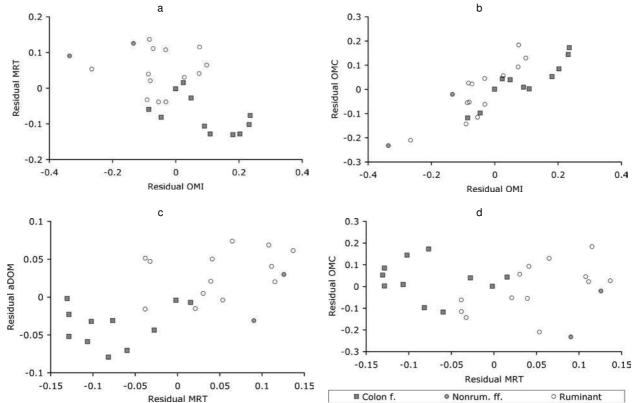


Fig. 1. Relationships between body mass residuals of different parameters of digestive
physiology (organic matter intake OMI in kg d⁻¹; particle mean retention time MRT in h;
organic matter gut contents OMC in kg, apparent organic matter digestibility aDOM in %) in
large mammalian herbivores of different digestion types (colon fermenters, nonruminant
foregut fermenters, ruminants) in the dataset of Foose (1982). a) OMI and MRT; b) OMI and
OMC; c) MRT and aDOM; d) MRT and OMC. For statistics, see Table 2.

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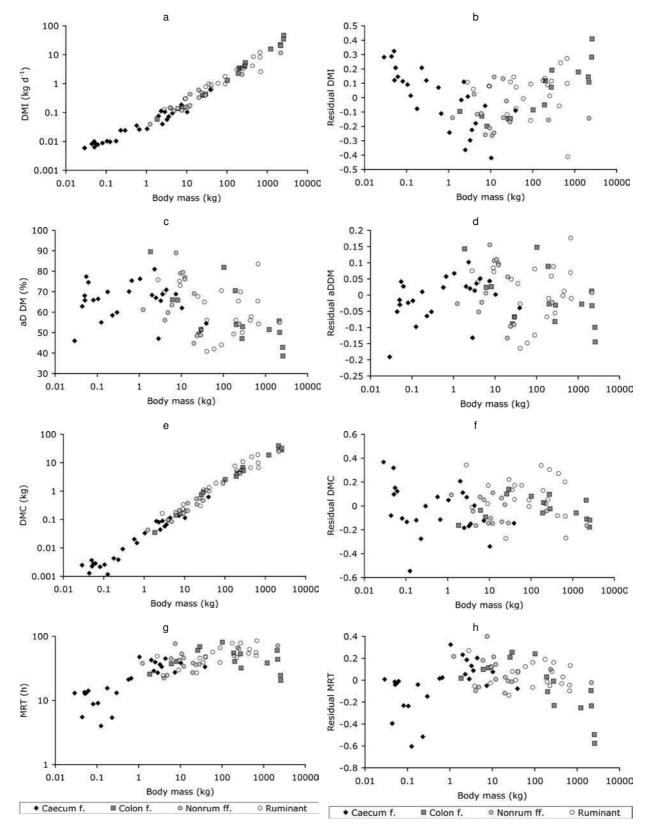
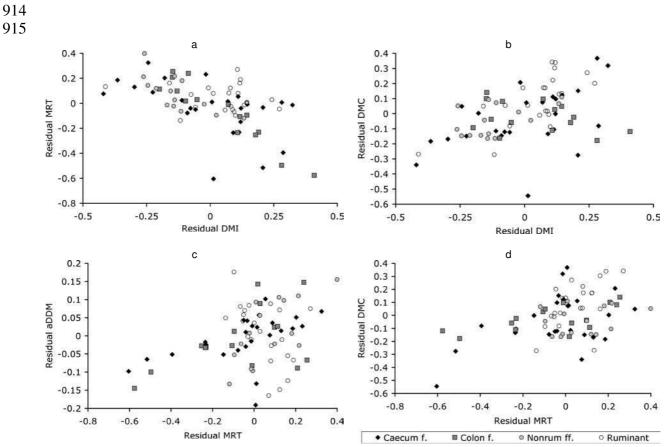


Figure 2. Scaling relationships between body mass (BM) and parameters of digestive physiology (left side) and
plots of body mass-residuals of these parameters against body mass to visualize patterns of quadratic scaling
(curvatures) (right side) in mammalian herbivores of different digestion types (caecum fermenters, colon
fermenters, nonruminant foregut fermenters, ruminants). a,b) dry matter intake (DMI); c,d) apparent digestibility
of dry matter (aD DM); e,f) dry matter gut contents (DMC); g,h) particle mean retention time (MRT). For
statistics, see Tables 3-6.



916 Fig. 3. Relationships between body mass residuals of different parameters of digestive 917 physiology (dry matter intake DMI in kg d⁻¹; particle mean retention time MRT in h; dry 918 matter gut contents DMC in kg, apparent dry matter digestibility aDDM in %) in mammalian 919 herbivores of different digestion types (colon fermenters, nonruminant foregut fermenters,

920 ruminants) in the dataset of this study. a) DMI and MRT; b) DMI and DMC; c) MRT and

921 922 aDDM; d) MRT and DMC. For statistics, see Table 7.

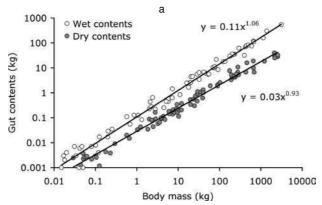


Figure 4. Scaling of wet gut contents (open symbols, Clauss et al., 2007a) and dry matter gut
 contents (closed symbols, this study) with body mass. Note the increasing difference between

926 the two with increasing body mass.



Assessing the Jarman-Bell-principle: scaling of intake, digestibility, retention time and gut capacity with body mass in mammalian herbivores

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Supplements

Table 1. Dataset from Foose (1982)

Gastrointestinal anatomy (GIT-Anatomy), mean of body mass (BM), organic matter intake (OMI), mean retention time of particle markers (different markers) through the whole gastrointestinal tract (MRT), apparent digestibility of organic matter (aD OM) and calculated mass of organic matter gastrointestinal tract content (OMC) of 26 herbivorous mammalian species for which all relevant data were measured on grass and lucerne hay diets

Species	GIT-Anatomy	BM (kg)	OM (kg/d)	MRT (h)	aD OM (%)	OMC (kg)
Tapirus terrestris	2	140	2.239	46	49.0	3.24
Equus asinus	2	183	5.747	45	53.8	7.86
Boselaphus tragocamelus	4	193	2.848	59	60.8	4.87
Cervus duvaucelii	4	193	2.822	50	64.5	3.98
Hexaprotodon liberiensis	3	200	1.649	67	53.9	3.34
Oryx gazella	4	204	4.318	71	60.7	8.84
Kobus ellipsiprymnus	4	204	3.859	58	58.6	6.59
Tapirus indicus	2	218	3.432	45	48.3	4.88
Equus zebra	2	265	7.553	44	54.0	9.98
Equus burchellii	2	272	7.200	41	55.2	8.91
Cervus elaphus	4	284	3.874	58	56.2	6.73
Syncerus caffer	4	291	5.954	64	69.0	10.44
Equus grevyi	2	353	8.347	42	58.1	10.24
Taurotragus oryx	4	431	5.959	52	56.5	9.27
Bison bison	4	499	6.651	74	68.8	13.36
Camelus bactrianus	4	544	6.308	79	67.7	13.73
Camelus dromedarius	4	544	6.477	75	64.5	13.62
Bos frontalis	4	765	8.728	54	66.5	13.11
Giraffa camelopardalis	4	1153	7.321	69	58.9	14.74
Bubalus bubalis	4	1270	17.232	67	66.8	32.03
Diceros bicornis	2	1285	16.419	57	53.8	28.62
Ceratotherium simum	2	1752	18.550	62	59.2	33.73
Rhinoceros unicornis	2	1852	20.468	65	58.9	38.96
Hippopotamus amphibius	3	2240	16.443	85	64.3	39.28
Loxodonta africana	2	2873	34.729	48	53.4	50.72
Elephas maximus	2	3034	34.748	51	52.6	53.86

GIT-Anatomy = gastrointestinal tract anatomy: 2 = colon fermenter, 3 = nonruminant foregut fermenter, 4 = ruminant foregut fermenter

Table 2. Data collection of this study

Gastrointestinal anatomy (GIT-Anatomy), number of offspring per litter (Nos offspring) mean of body mass (BM), dry matter intake (DMI), mean retention time of particle markers (different markers) through the whole gastrointestinal tract (MRT), apparent digestibility of dry matter (aD DM) and calculated mass of dry matter gastrointestinal tract content (DMC) of 77 herbivorous mammalian species for which all relevant data were measured in individual studies

Species	GIT- Anatomy	Nos offspring	BM (kg)	DMI (kg/d)	MRT (h)	aD DM (%)	DMC (kg)	Source
Mus musculus	1	>1	0.03	0.006	13.0	46.0	0.002	(Karasov et al., 1986)
Lasiopodomys brandtii	1	>1	0.04	0.008	5.5	62.9	0.001	(Pei et al., 2001a)
Meriones unguiculatus	1	>1	0.05	0.006	12.8	65.8	0.002	(Pei et al., 2001b)
Microtus ochrogaster	1	>1	0.05	0.010	13.5	68.1	0.004	(Hammond and Wunder, 1991)
Microtus townsendii	1	>1	0.05	0.008	13.1	77.4	0.003	(Hume et al., 1993)
Tamias amoenus	1	>1	0.06	0.008	14.1	74.6	0.003	(Hume et al., 1993)
Phyllotis darwini	1	>1	0.08	0.009	8.8	65.9	0.002	(Sakaguchi and Ohmura, 1992)
Cricetus cricetus	1	>1	0.11	0.011	9.1	66.5	0.003	(Sakaguchi et al., 1987)
Neotoma lepida	1	>1	0.13	0.010	4.0	55.0	0.001	(Karasov et al., 1986)
Octodon degus	1	>1	0.18	0.010	15.5	69.9	0.004	(Sakaguchi and Ohmura, 1992)
Arvicola terrestris	1	>1	0.23	0.024	5.4	58.5	0.004	(Woodall, 1989)
Rattus norvegicus	1	>1	0.30	0.024	13.1	59.9	0.009	(Sakaguchi et al., 1987)
Cavia porcellus	1	>1	0.57	0.036	21.1	70.1	0.020	(Sakaguchi et al., 1987; Sakaguchi and Nabata, 1992; Sakaguchi et al., 1992b; Sakaguchi and Ohmura, 1992; Franz et al., 2011)
Spermophilus columbianus	1	>1	0.66	0.026	22.1	75.4	0.015	(Hume et al., 1993)
Hapalemur griseus	1	1	1.04	0.027	47.5	76.3	0.033	(Campbell et al., 2004)
Oryctolagus cuniculus	1	>1	2.01	0.079	42.6	68.4	0.088	(Sakaguchi et al., 1987; Sakaguchi and Hume, 1990; Sakaguchi et al., 1992a; Franz et al., 2011)
Marmota caligata	1	>1	2.31	0.112	28.9	81.0	0.080	(Hume et al., 1993)
Trichosurus vulpecula	1	1	2.50	0.040	39.6	67.0	0.044	(Sakaguchi and Hume, 1990)
Lagostomus maximus	1	>1	2.86	0.105	27.0	47.0	0.090	(Besselmann, 2005; Clauss et al., 2007a)
Propithecus tattersalli	1	1	3.24	0.057	36.3	65.6	0.058	(Campbell et al., 1999)
Propithecus verreauxi	1	1	3.58	0.072	33.5	68.8	0.066	(Campbell et al., 1999; Campbell et al., 2004)
Myocastor coypus	1	>1	4.40	0.094	45.0	70.9	0.114	(Sakaguchi and Nabata, 1992)
Dolichotis patagonum	1	>1	7.40	0.185	27.3	68.8	0.138	(Sakaguchi et al., 1992b)
Erethizon dorsatum	1	1	10.3	0.103	38.4	62.0	0.114	(Felicetti et al., 2000)
Hydrochaeris hydrochaeris	1	>1	39.2	0.606	33.3	54.5	0.620	(A. Schwarm, S. Ortmann, M. Clauss pers. obs. 2010)

Pithecia pithecia	2	1	1.85	0.059	25.6	89.5	0.035	(Norconk et al., 2002)
Alouatta pigra	2	1	6.15	0.137	37.3	66.1	0.142	(Edwards and Ullrey, 1999)
Alouatta seniculus	2	1	8.18	0.144	40.4	65.9	0.163	(Edwards and Ullrey, 1999)
Lasiorhinus latifrons	2	1	26.2	0.394	60.5	49.2	0.750	(Barboza, 1993)
Vombatus ursinus	2	1	29.5	0.434	68.5	51.6	0.918	(Barboza, 1993)
Gorilla gorilla	2	1	102	1.267	80.6	81.9	2.560	(Remis and Dierenfeld, 2004)
Tapirus terrestris	2	1	192	2.247	54.7	70.5	3.320	(Clauss et al., 2010)
Equus asinus	2	1	202	3.445	40.5	53.9	4.250	(Pearson and Merritt, 1991; Pearson et al., 2001, 2006)
Tapirus indicus	2	1	278	3.945	53.1	47.1	6.720	(Clauss et al., 2010)
Equus caballus	2	1	287	5.416	32.0	52.9	5.250	(Pearson and Merritt, 1991; Pagan et al., 1998; Pearson et al., 2001; Moore-Colyer et al., 2003; Pearson et 2006)
Diceros bicornis	2	1	1222	15.700	38.3	51.4	18.614	(Clauss et al., 2005a; Clauss et al., 2006; Steuer et al., 2010)
Rhinoceros unicornis	2	1	2125	22.065	60.1	55.7	39.935	(Clauss et al., 2005b)
Ceratotherium simum	2	1	2175	20.781	43.8	50.1	28.286	(Steuer et al., 2010)
Elephas maximus	2	1	2547	33.874	24.5	42.8	27.998	(Hackenberger, 1987)
Loxodonta africana	2	1	2583	47.207	20.5	38.6	32.515	(Hackenberger, 1987)
Lagorchestes hirsutus	3	1	1.23	0.040	38.0	61.3	0.043	(Bridie et al., 1994)
Thylogale thetis	3	1	4.05	0.141	22.4	56.1	0.095	(Dellow, 1982; Dellow and Hume, 1982)
Macropus eugenii	3	1	4.80	0.117	24.8	59.8	0.084	(Dellow, 1982; Dellow and Hume, 1982)
Trachypithecus auratus	3	1	6.00	0.140	45.0	63.5	0.179	(Nijboer et al., 2007)
Colobus angolensis	3	1	7.50	0.118	77.0	89.0	0.210	(Schwarm et al., 2009a)
Trachypithecus johnii	3	1	9.50	0.157	42.0	79.0	0.166	(Comizzoli et al., 1997; Schwarm et al., 2009a)
Colobus guereza	3	1	10.8	0.154	53.1	79.5	0.206	(Edwards and Ullrey, 1999)
Colobus polykomos	3	1	12	0.174	46.0	77.0	0.205	(Schwarm et al., 2009a)
Pygathrix nemaeus	3	1	12.1	0.429	33.8	76.2	0.374	(Edwards and Ullrey, 1999)
Macropus robustus	3	1	19.8	0.608	27.2	44.7	0.534	(Freudenberger and Hume, 1992)
Pecari tajacu	3	>1	20.1	0.324	38.2	69.1	0.343	(Schwarm et al., 2009a)
Macropus giganteus	3	1	23.6	0.504	36.2	48.4	0.555	(Forbes and Tribe, 1970; Dellow, 1982; Dellow and Hume, 1982)
Macropus rufus	3	1	28.7	0.468	36.7	48.8	0.467	(Foot and Romberg, 1965; McIntosh, 1966; Forbes and Tribe, 1970; Dellow, 1982; Dellow and Hume, 19 Munn and Dawson, 2006; Schwarm et al., 2009a)
Hexaprotodon liberiensis	3	1	238	2.067	66.5	54.3	4.215	(Clauss et al., 2004)
Hippopotamus amphibius	3	1	2175	11.733	71.4	55.0	25.011	(Clauss et al., 2004)
Tragulus napu	4	1	2.8	0.129	49.0	75.8	0.164	(Conklin and Dierenfeld, 1994)

Cephalophus monticola	4	1	3.85	0.146	24.4	69.6	0.098	(Luginbuhl et al., 1990)
Pudu puda	4	1	9.1	0.299	29.9	75.2	0.232	(Conklin-Brittain and Dierenfeld, 1996)
Cephalophus maxwellii	4	1	9.4	0.305	42.2	73.0	0.341	(Conklin-Brittain and Dierenfeld, 1996)
Capreolus capreolus	4	>1	24.7	0.405	27.0	67.7	0.302	(Holand, 1994)
Capra hircus	4	1	29.7	0.777	45.9	51.3	1.117	(Udén et al., 1982; Udén and Van Soest, 1982; Freudenberger and Hume, 1992; Kennedy et al., 1992)
Capra ibex	4	1	35.4	0.989	39.3	65.0	1.074	(Gross et al., 1996)
Odocoileus virginianus	4	>1	40.3	0.648	47.0	40.9	1.011	(Barnes et al., 1992)
Ovis ammon	4	1	40.3	0.914	48.4	56.1	1.325	(Foot and Romberg, 1965; McIntosh, 1966; Forbes and Tribe, 1970; Dellow, 1982; Dellow and Hume, 1982; Udén et al., 1982; Udén and Van Soest, 1982; Luginbuhl et al., 1990; Kennedy et al., 1992; Pearson et al., 2006)
Odocoileus hemionus	4	>1	60	1.028	56.7	42.0	1.918	(Baker and Hobbs, 1987)
Bubalus depressicornis	4	1	90	1.767	39.0	70.5	1.858	(Flores-Miyamoto et al., 2005)
Ovis canadensis	4	1	90	0.984	66.0	44.0	2.111	(Baker and Hobbs, 1987)
Bos grunniens	4	1	176	3.122	78.2	49.3	7.663	(Schaefer et al., 1978)
Cervus elaphus	4	1	196	3.763	49.7	56.1	5.226	(Baker and Hobbs, 1987; Jiang and Hudson, 1996)
Okapia johnstoni	4	1	227	3.531	46.8	65.4	4.626	(Hummel et al., 2005; Hummel et al., 2006)
Ovibos moschatus	4	1	254	3.223	64.2	69.9	5.623	(Adamczewski et al., 1993; Barboza et al., 2006)
Bison bison	4	1	279	4.410	78.8	50.1	10.852	(Schaefer et al., 1978)
Bos javanicus	4	1	432	4.654	54.2	56.0	6.711	(Schwarm et al., 2008, 2009b)
Bos taurus	4	1	460	7.357	62.8	57.8	16.052	(Schaefer et al., 1978; Colucci et al., 1982; Udén et al., 1982; Udén and Van Soest, 1982; Mathers et al., 1989; Luginbuhl et al., 1994; Burns et al., 1997; Whetsell et al., 2004; Pearson et al., 2006)
Bubalus bubalis	4	1	656	12.120	56.0	65.4	19.037	(Mathers et al., 1989)
Giraffa camelopardalis	4	1	665	8.224	49.8	83.5	9.942	(Schaub, 2005)
Camelus bactrianus	4	1	687	2.601	85.2	54.3	6.724	(Cahill and McBride, 1995)

GIT-Anatomy = gastrointestinal tract anatomy: 1 = caecum fermenter, 2 = colon fermenter, 3 = nonruminant foregut fermenter, 4 = ruminant foregut fermenter; BM = body mass; DMI = dry matter intake; MRT = mean retention time of particle markers (different markers) through the whole gastrointestinal tract; aD DM = apparent dry matter digestibility; DMC= mass of dry matter gastrointestinal tract content calculated according to Holleman and White (1989; for equations see the main text)

Table 3. Data collection of wet gut contentsThis dataset is given as the supplement to Clauss et al. (2007b).

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