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1           **Full Title: Introduced megafauna are rewilding the Anthropocene**

2                                   **Short title: Anthropocene megafauna**

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23 **ABSTRACT**

24 Large herbivorous mammals, already greatly reduced by the late-Pleistocene extinctions,  
25 continue to be threatened with decline. However, many herbivorous megafauna (body  
26 mass  $\geq$  100 kg) have populations outside their native ranges. We evaluate the  
27 distribution, diversity and threat status of introduced terrestrial megafauna worldwide and  
28 their contribution towards lost Pleistocene species richness. Of 76 megafauna species, 22  
29 (~29%) have introduced populations; of these ten (45%) are threatened or extinct in their  
30 native ranges. Introductions have increased megafauna species richness by between 10%  
31 (Africa) and 100% (Australia). Furthermore, between 15% (Asia) and 67% (Australia) of  
32 extinct species richness, from the late Pleistocene to today, have been numerically  
33 replaced by introduced megafauna. Much remains unknown about the ecology of  
34 introduced herbivores, but evidence suggests that these populations are rewilding modern  
35 ecosystems. We propose that attitudes towards introduced megafauna should allow for  
36 broader research and management goals.

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46 **INTRODUCTION**

47 Terrestrial herbivorous megafauna are undergoing severe declines around the  
48 world. Of 74 extant large terrestrial herbivorous mammal species with body masses  $\geq 100$   
49 kg, 44 (~60%) are threatened with extinction (Ripple et al. 2015). The decline of this  
50 functional group began 10,000-50,000 years ago, most likely due to overhunting by  
51 humans during the late Pleistocene (Barnosky et al. 2004, Bartlett et al. 2015).

52 Large ( $\geq 100$  kg) herbivorous megafauna (henceforth 'megafauna') perform  
53 distinct roles that contribute to the functioning of ecological systems. Megafauna  
54 consume fibrous vegetation, which can benefit smaller herbivores, reduce fire risk,  
55 accelerate rates of nutrient cycling by orders of magnitude, and shift plant community  
56 structure by facilitating coexistence between different plant functional types. Due to their  
57 large size, these organisms cause physical disturbance and disperse large seeds and  
58 nutrients great distances (Ripple et al. 2015). The considerable loss of this functionality at  
59 the end of the Pleistocene had dramatic effects on plant community structure, fire  
60 regimes, nutrient and mineral cycling across landscapes, and community assembly  
61 (Bakker et al. 2016a, Gill et al. 2009, Doughty et al. 2016a, Doughty et al. 2016b,  
62 Doughty et al. 2016c, Ripple & Van Valkenburgh 2010, Smith et al. 2015, Malhi et al.  
63 2016). Modern declines have similar consequences for terrestrial ecosystems and  
64 community dynamics (Ripple et al. 2015) and have led to broad international calls for  
65 immediate action to conserve the world's remaining mammalian megafauna (Ripple et al.  
66 2016, Ripple et al. 2017).

67 Less well considered is the role of megafauna introductions on their conservation  
68 and on ecosystem function. Since the advent of the Anthropocene, particularly in the past

69 200 years, megafauna have been moved to new regions and between continents.  
70 Introductions of megafauna worldwide may have inadvertently provided refuge for  
71 threatened megafauna, increased regional large herbivore species richness, and restored  
72 or added ecological functions. Acknowledgement of this possibility is being fostered by  
73 the burgeoning concept of ‘rewilding,’ which includes efforts to proactively introduce  
74 species in order to provide refuge and to restore lost ecological processes (Donlan et al.  
75 2006, Svenning et al. 2016). However, much remains unknown about the contribution of  
76 already introduced populations to global conservation goals.

77         Given that introduced populations are often unwanted and considered components  
78 of anthropogenic harm, the existence of populations that are simultaneously introduced  
79 and threatened or extinct in their native ranges has been highlighted as a conservation  
80 paradox (Marchetti and Engstrom 2015). Indeed, the considerable redistribution of biota  
81 that characterizes the Anthropocene may be a countercurrent to the extinction crisis by  
82 providing refuge and new opportunities for threatened species (Wallach et al. 2015).  
83 However, comprehensive analyses of the interaction between the processes of extinction  
84 and redistribution have not been conducted.

85         To assess the potential conservation values of introduced megafauna we compiled  
86 current information on their threat statuses and population trends in their native ranges,  
87 their relative population sizes in and out of their native ranges, and their functional roles.  
88 To understand how introduced megafauna have potentially rewilded the world, we  
89 assessed the contribution of introduced megafauna to continental assemblages, and the  
90 contribution of introduced megafauna to Anthropocene richness relative to the Holocene  
91 and Pleistocene epochs.

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## 93 **METHODS**

94           We searched for introduced populations of herbivorous megafauna (mammals  
95 only) with body masses  $\geq 100$  kg based on Ripple et al. (2015) using Long (2003) and  
96 supplemented with online searches (Google Scholar and Google) using the terms “feral”,  
97 “introduced”, “invasive”, “exotic” and “non-native”. We used grey literature (e.g.  
98 government reports) and journalism sources (e.g. The New York Times) alongside peer-  
99 reviewed literature to identify megafauna populations outside their native ranges. Data  
100 collection concluded in July, 2017. While some native megafauna populations live in  
101 fenced and managed conditions (e.g. Kruger National Park), only free-roaming wild  
102 introduced populations were included because it was not clear if fenced/managed  
103 introduced populations are ecologically viable in their new homes.

104           To understand to what extent introduced megafauna represent the taxonomic  
105 diversity of the world’s remaining megafauna, we calculated the number of large  
106 herbivore families represented by introduced species, the number of genera of each  
107 family represented by introduced species, and the percentage of species with introduced  
108 populations within each taxonomic family.

109           To determine the potential conservation value of introduced megafauna as refuge  
110 populations, we compiled IUCN (2017) Red List threat statuses and trends in each  
111 species’ historic native ranges and the proportion of each population that is currently  
112 outside of its native range (Supplementary material Appendix 2, Table A1). Wild post-  
113 domestic species were assigned the threat status of their pre-domestic ancestor. For  
114 example, introduced wild dromedary camels (*Camelus dromedarius*) originate from the

115 domesticated form of an extinct camel species (possibly *C. thomasi*), and were therefore  
116 considered extinct in the wild in their native range.

117 To understand to what geographic extent introduced megafauna have rewilded the  
118 world, we calculated megafauna species richness by Taxonomic Databases Working  
119 Group level 3 countries (henceforth TDWG), which are bio-geographic units defined by  
120 political (nation, state, province, or district) boundaries at a biologically relevant scale  
121 (Brummitt 2001). Inter- and intra-continental introductions were included in this  
122 comparison. The distributions of introduced megafauna were determined from literature  
123 and Google searches (Supplementary material Appendix 1, Data A1). Geographic ranges  
124 for native megafauna were downloaded from the IUCN (2017) Red List. The percentage  
125 of each TDWG country's megafauna assemblage that is introduced was calculated and  
126 compared between continents to understand how introductions have altered continental  
127 megafauna assemblages.

128 We assessed how Anthropocene megafauna richness compares to those of past  
129 geological epochs. For each continent, we compared megafauna species richness and  
130 conservation status between the late Pleistocene (50,000 – 10,000 BP), Holocene  
131 (<10,000 BP), and Anthropocene (past ~ 200 years) epochs. Only inter-continental  
132 introduced megafauna were included.

133 Pleistocene species were classified as 'extinct', 'extirpated' or 'survived' based on  
134 their fate through the late-Pleistocene extinction. Pleistocene megafauna presence was  
135 based on Sandom et al. (2014) and body masses ( $\geq 100\text{kg}$ ) were confirmed through  
136 literature searches. The Holocene included species from the end of the Pleistocene until  
137 the Anthropocene. Holocene species included 'survived' taxa, natural immigrants, and

138 species that went extinct during the Holocene (e.g. aurochs *Bos primigenius* and  
139 dromedary camel), while Anthropocene genera included ‘survived’, ‘survived,  
140 threatened’, ‘introduced’, and ‘introduced, threatened’ species, reflecting their current  
141 IUCN (2017) threat statuses (Supplementary material Appendix 3, Table A2).

142 To describe the range of functional traits of introduced megafauna, we reviewed  
143 their average body masses, habitat types, dietary types (grazer, browser, or intermediate),  
144 and other unique traits using the IUCN (2017) and published literature.

145

## 146 **RESULTS**

147 Twenty-two (32%) of the 76 extant megafauna species have established wild  
148 populations outside their native ranges (Supplementary material Appendix 2, Table A1).  
149 Sixteen are inter-continental introductions, two are intra-regional but overcame oceanic  
150 barriers, and four are intra-continental. By including post-domesticates of extinct  
151 heritage, an additional two species (the dromedary camel and cattle, *Bos taurus*) are  
152 added to the 74 remaining native megafauna. Six additional species were excluded from  
153 analysis: three species because they appear to be confined to game ranches, one because  
154 introduced populations are described as semi-wild, and two because of uncertain  
155 taxonomic relation to already included species.

156 Six (55%) of the eleven families containing megafauna species have established  
157 populations outside their native ranges. Introduced species represent between 29%  
158 (Equidae) and 56% (Cervidae) of the megafauna species within their families (Fig. 1).  
159 Likewise, introduced populations represent between 50% (Camelidae) and 100%  
160 (Equidae) of the megafauna genera within their families.



161           Of the 22 species with introduced populations, ten (45%) are threatened or extinct  
162 in their native ranges (Fig. 2). This includes three (~14%) Vulnerable non-domesticated  
163 species, four (18%) post-domestics whose progenitors are Endangered, two (9%) post-  
164 domestic species whose wild progenitors are Extinct, and one (~5%) post-domestic  
165 whose progenitor is Critically Endangered. All seven post-domestic species are extinct or  
166 threatened in their native ranges. Of the remaining twelve introduced megafauna, three  
167 (14%) are Near Threatened, and nine are ranked as Least Concern in their native ranges,  
168 of which 66% have stable population trends, 22% are increasing, and 11% are declining  
169 (Fig. 2). Of the 20 introduced species with surviving native populations, ten (50%) are  
170 declining in their native ranges, seven (35%) are stable, and three (15%) are increasing  
171 (Supplementary material Appendix 2, Table A1). In all, 59% of introduced megafauna  
172 are threatened or declining in their native ranges (Fig. 2).

173           On average, over 38% (ranging between <1% and 100%) of megafauna  
174 populations are outside of their native ranges. Whereas two species have relatively small  
175 (possibly ~100 individuals) populations outside their native ranges (hippopotamus,  
176 *Hippopotamus amphibius*, and Asian elephant, *Elephas maximus*), twelve populations are  
177 estimated in the thousands and up to over 1 million individuals (Fig. 3, Supplementary  
178 material Appendix 2, Table A1).

179           By including introduced megafauna, the worldwide distribution of megafauna  
180 species richness increases significantly (Fig. 4). Introduced megafauna have substantially  
181 increased continental megafauna richness and TDWG-country-scale species richness  
182 within each continent: 62% of South American (mean = 37%, SD = 34%), 57% of North  
183 American (TDWG: mean = 24%, SD = 37%), 33% of European (mean = 36%, SD =

184 33%), 11% of Asian (mean = 17%, SD = 34%), and 11% of African (mean = 10%, SD =  
185 27%) megafauna are introduced. Introduced megafauna comprise at least 75% of the  
186 megafauna assemblages of 56 of the 369 (15%) TDWG countries.

187 Strikingly, the entire continental megafauna assemblage of Australia is composed  
188 of introduced species. Australia lost all megafauna species during the Pleistocene  
189 extinctions, yet has become home to eight introduced species in the Anthropocene,  
190 including the Endangered Banteng (*Bos javanicus*), the world's only population of wild  
191 dromedary camel, the Vulnerable sambar deer (*Rusa unicolor*), and the water buffalo  
192 (*Bubalus bubalis*), the descendant of the Endangered water buffalo (*B. arnee*). Wild  
193 donkeys (*Equus asinus*), whose progenitor, the African wild ass (*E. africanus*) is  
194 Critically Endangered, and Endangered horses (*E. ferus caballus*), have also found refuge  
195 in Australia, as well as in North America, South America, and Europe.

196 Late Pleistocene losses of megafauna species (100% for Australia, 89% for South  
197 America, 89% for North America, 53% for Europe, 41% for Asia, and 27% for Africa)  
198 and Holocene losses (14% for Europe, 5% for Asia, and 3% for Africa) were substantial.  
199 Following the Pleistocene, North American species richness increased from 4 to 6 due to  
200 immigration of wapiti (*Cervus canadensis*) and moose (*Alces alces*) from Eurasia  
201 concurrent with the arrival of the first humans to the continent (Meiri *et al.* 2014,  
202 Hundertmark *et al.*, 2002). Reductions in species richness on all continents since the  
203 Pleistocene have been counteracted by gains from introduced megafauna in the  
204 Anthropocene, so that there are currently more megafauna species per continent than at  
205 the end of the Holocene. Introduced megafauna have numerically replaced extinct species

206 richness in Australia by 67%, in South America by 21%, in North America by 26%, in  
207 Europe by 33%, in Asia by 15%, and in Africa by 31% (Fig. 5, Table 1).

208         Megafauna are likely to have significant functional roles in their introduced  
209 ranges. Their average body masses ranges from 109 to 3,270 kg (median = 256 kg, mean  
210 = 526 kg, SD = 697 kg) (Table 2), which is representative of the native megafauna body  
211 mass distribution ranging from 100 to 3,825 kg (median = 238 kg, mean = 496 kg, SD =  
212 666 kg). Introduced megafauna are primarily grazers (45% of species) or intermediate  
213 grazers and browsers (41% of species), and three species (14%) are primarily browsers  
214 (Table 1). Introduced megafauna are adapted for habitats ranging from Arctic tundra  
215 (muskox *Ovibos moschatus*) to tropical forest (sambar deer) and deserts (dromedary  
216 camels) (Table 1). Although there is little known about the specific ecological  
217 functionalities of several introduced megafauna, many introduced species are known for  
218 unique traits, such as the ability to drink brackish water and consume halophytic plants  
219 (dromedary camel) or to survive without surface water (gemsbok *Oryx gazella*) (Table 2).

220

## 221 **DISCUSSION**

222         Introduced megafauna represent a significant proportion of the remaining  
223 taxonomic diversity of their functional group and are themselves significantly threatened  
224 in their historic native ranges. This raises the question of how to assign conservation  
225 value in an era of extinction and redistribution. Conservation biology is a field driven by  
226 a plurality of values, which offer various visions at different scales and times (Sandbrook  
227 et al. 2011). Many current schools of thought prioritize the conservation of species  
228 considered to be native at the local and regional scale. However, given the ongoing global

229 extinction process, more research and dialogue is needed to understand when these values  
230 may undermine other conservation goals and values.

231         While many introduced populations were formerly domesticated, they may still  
232 effectively represent their wild relatives. Introduced populations of Endangered banteng  
233 in northern Australia have maintained high genetic fidelity to their pre-domestic  
234 ancestors (Bradshaw et al. 2005). Likewise, domesticated horses retain a substantial  
235 component of the genetic diversity of extinct Holarctic horse lineages (Lippold et al.  
236 2011). Given that the closest wild relatives of all six post-domestic megafauna are  
237 Endangered or extinct, it appears that domestication has provided a crucial bridge for  
238 certain species from the pre-pastoral wild landscapes of the early Holocene to the post-  
239 industrial wild landscapes of the Anthropocene.

240         Evolutionary and ecological change has also been witnessed in post-domestic  
241 populations. Wild goats (*Capra aegragus*) on Aldabra Atoll regularly drink saltwater  
242 when freshwater is absent (Burke 1990). Wild sheep (*Ovis aries*) show higher resistance  
243 to local parasites than sympatric domestic sheep. Wild Ossabaw island pigs (*Sus scrofa*)  
244 have unique lipid structures (Van Vuren and Hedrick 1989). Wild cattle in Mexico do not  
245 linger in riparian areas like their sympatric domestic cousins due to altered predation  
246 threats (Hernandez et al. 1999). Native Torresian crows (*Corvus orru*) appear to have  
247 developed a mutualistic grooming behavior on introduced banteng in Australia  
248 (Bradshaw and White 2006).

249         Like all herbivores, introduced megafauna can exert strong grazing or browsing  
250 pressure to the detriment of other species, most notably where apex predators are  
251 extirpated or continue to be persecuted (Wallach et al. 2010). Unfortunately, much of the

252 research to document these effects has ignored the ecological context of predator control,  
253 which is to ignore an important explanatory variable for the density-dependent effects of  
254 all herbivores. Indeed, wild horses in the United States may be limited by mountain lions  
255 (Turner and Morrison 2001) and dingoes appear to suppress populations of wild donkeys  
256 in Australia (Wallach et al. 2010). The potential to influence the ecologies of introduced  
257 megafauna by protecting or restoring large predators is an important topic for further  
258 research.

259         In the Pleistocene, the ecological influences of herbivorous megafauna on  
260 disturbance regimes, seed dispersal, nutrient cycling, and community structure were  
261 ubiquitous. Introduced megafauna have potentially augmented this lost functional and  
262 taxonomic diversity across most continents, particularly in those regions most depleted:  
263 Australia, North America, and South America (Fig. 4); Asia and Africa have retained  
264 many Pleistocene megafauna and have fewer introduced species. Several of these  
265 introductions restore taxonomic analogues to extinct Pleistocene species. For example,  
266 introduced donkeys are morphologically similar to congeneric extinct North American  
267 and South American stilt-legged horses, and the modern wild horse is the same species as  
268 the horse of the Holarctic Pleistocene (Weinstock et al. 2005).

269         The late Pleistocene extinctions in Australia included all megafauna and many  
270 browsing herbivores, the loss of which appears to have led to increased fire frequency  
271 and altered plant community structure (Miller et al. 2005, Rule et al. 2012). Introduced  
272 megafauna, especially browsers such as dromedary camels, may reverse these ecological  
273 state shifts. However, determining how introductions of taxonomically dissimilar species  
274 restore or add new functionalities within insular ecosystems (there are no surviving

275 taxonomic analogues to Australia's Pleistocene marsupial megafauna) requires further  
276 research into the relative importance of co-evolutionary history versus ecological context  
277 in determining species coexistence and ecosystem function (Wallach et al. 2015).

278         Introduced megafauna vary in body mass considerably, which influences their  
279 ability to open thickets and digest coarse fibrous vegetation and thus their relation to  
280 plant communities and other herbivores. Introduced megafauna also possess unique  
281 functional adaptations that may be of ecological significance in their new ranges. For  
282 example, introduced camels are capable of ingesting brackish water and consuming  
283 halophytic plants (Root-Bernstein & Svenning 2016), which in conjunction with their  
284 large home ranges (Spencer et al. 2012) may contribute to the megafaunal redistribution  
285 of terrestrial salts (Doughty et al. 2016a). Likewise, the ability of gemsbok (*Oryx gazella*)  
286 to survive without surface water (Hamilton et al. 1977) likely allows it to occupy novel  
287 niches in the North American deserts in which it now lives.

288         There is substantial and growing evidence that introduced species can perform  
289 significant and desirable ecological roles (Schlaepfer et al. 2011). Bighorn sheep forage  
290 more efficiently, with less time invested in vigilance behaviors in mixed herds with  
291 introduced wild horses (Coates and Schemnitz 1994). Giant tortoises introduced onto  
292 oceanic islands as substitutes for extinct species are dispersing large-seeded endemic  
293 plants and shaping plant communities through grazing (Hansen et al. 2010). Intentional  
294 introductions of horses and cattle in the Oostvaardersplassen nature reserve in the  
295 Netherlands have created Pleistocene-like savanna conditions in a temperate deciduous  
296 forest environment (Vera 2009). In North America and Australia, the drying and  
297 constriction of desert springs and the extinction of several endemic fish populations was

298 linked to the removal of wild introduced megafauna whose grazing appeared to maintain  
299 open-water habitat (Kodric-Brown and Brown 2007).

300         Likewise, our own ongoing research is yielding similarly surprising observations.  
301 For example, in the Sonoran Desert of North America, wild donkeys ('burros', *E. asinus*)  
302 dig groundwater wells of more than a meter in depth (Supplementary material Appendix  
303 4, Movie A1). These wells are common wherever groundwater approaches the surface,  
304 have been recorded in use by more than thirty mammal and bird species, and in certain  
305 conditions become nurseries for riparian trees (Fig. 6). It is possible that by creating new  
306 water sources across the landscape, maintaining access to receding water-tables during  
307 droughts, and providing conditions ideal for the germination of riparian trees, wild  
308 donkeys play a facilitative role, one that may improve the resilience of these arid  
309 ecosystems to climate change. Furthermore, given the ubiquity of taxa whose  
310 contemporaries dig wells, such as Proboscideans (Ramey et al. 2013) and other equids  
311 (Feh et al. 2002) in the North American Pleistocene, it is likely that introduced donkeys  
312 have restored a functionality lost from these landscapes.

313         Unfortunately, little more is known about the ecological functions of megafauna  
314 outside their native ranges because the majority of studies are conducted on the premise  
315 that introduced species are harmful and should be suppressed or eradicated. Future  
316 research on the ecological functions of introduced megafauna, under varying ecological  
317 contexts (e.g. predator control, landscape connectivity), will be essential to understand  
318 the novel megafaunal communities of the Anthropocene.

319         Reassessing conservation attitudes towards introduced megafauna may find  
320 synergy with other conservation goals. Introduced megafauna are likely vulnerable to

321 similar threats as native megafauna as they require large tracts of land and may be  
322 vulnerable to exploitation. Valuing introduced megafauna as umbrella or flagship species  
323 in efforts to expand protected areas or establish movement corridors would contribute to  
324 important conservation goals. Broadening the range of wildlife valued and protected by  
325 conservation practitioners could also help form alliances with public advocates of  
326 introduced megafauna, who are often alienated by projects that treat these species as  
327 pests. Conflicts between these groups and conservation professionals erode trust and  
328 undermine conservation efficacy (Crowley et al. 2017), yet these groups are natural allies  
329 in their concern for the welfare and persistence of non-human life (Bruskotter et al.  
330 2017). It is likely that incorporating broader value systems towards these organisms  
331 would offer a range of practical benefits towards conservation objectives and could  
332 strengthen the diversity and inclusiveness of the conservation community.

333         The introduced megafauna of the world have restored species richness across  
334 many continents to levels approaching the Pleistocene, contribute fascinating and  
335 potentially important ecological functions, and are an important refuge for their  
336 functional group. We propose that further research and dialogue on how introduced  
337 megafauna interact with and without potential predators in the novel ecosystems of the  
338 Anthropocene will be essential in reconciling the concerns of local managers with global  
339 conservation efforts and will bring new attention to the emerging eco-evolutionary  
340 trajectories of these populations.

341

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602

### 603 **ACKNOWLEDGEMENTS**

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611 Table 1. Changes in megafauna species richness from the Pleistocene to the Anthropocene. In column 2, percent survived is the  
612 percent of megafauna to survive the late Pleistocene extinctions; in column 3, percent lost/gained is the percent change in Holocene  
613 species richness due to extinction/immigration during the Holocene; in column 4, percent replaced is the percent of all extinct  
614 megafauna richness (Pleistocene and Holocene) to be numerically replaced by introductions in the Anthropocene. \* indicates natural  
615 immigration from Eurasia to North America during the early Holocene.

Continent	Pleistocene species richness	Holocene species richness (percent survived)	Holocene extinctions/immigration (percent lost/gained)	Anthropocene richness (percent replaced)
Africa	44	32 (73%)	-1 (-3%)	35 (31%)
Asia	61	36 (59%)	-2 (-6%)	38 (14%)
Europe	15	7 (47%)	-1 (-14%)	9 (33%)
North America	35	4 (11%)	+2 (+33%)*	14 (26%)
South America	44	5 (11%)	0 (0%)	12 (18%)
Australia	12	0 (0%)	N/A	8 (67%)

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618 Table 2. Functional traits of introduced megafauna. ABM is average body mass (Jones et al. 2009); foraging type is 'B' is browser, 'G' is  
 619 grazer, and G/B are intermediate; habitats are derived from IUCN Redlist species accounts (give web address here).

Species	Common Name	ABM	Type	Habitat	Known or potential unique ecological functions
<i>Alces alces</i>	Moose	541	B	Woodlands, tundra, montane forests	Browse at heights up to 2m, affecting stand height and canopy composition (Pastor et al. 1988).
<i>Bison bison</i>	Bison	625	G	Grasslands, open forests	Create wallows that become ephemeral pools, serve as fire breaks, and increase landscape scale plant diversity (Knapp et al. 1999).
<i>Bos javanicus</i>	Banteng	636	G/B	Open dry forests	
<i>Bos taurus</i>	Cattle	613	G	Numerous	
<i>Boselaphus tragocamelus</i>	Nilgai	182	G/B	Open grasslands	Open trails in dense shrubland, capable of jumping 2.5m high-potentially sustaining seed/nutrient dispersal in fenced landscapes (Leslie 2008).
<i>Bubalus bubalis</i>	Water buffalo	919	G/B	Moist grasslands, marshes	Used for conservation grazing to maintain open water habitat for birds and fish (BBC News).
<i>Camelus dromedarius</i>	Dromedary camel	488	B	Desert scrub	Salt-tolerant (Root-Bernstein & Svenning 2016); large home ranges (Spencer et al. 2012), may redistribute sodium (Doughty et al. 2016a).
<i>Cervus elaphus</i>	Red deer	241	G/B	Generalist	
<i>Connochaetes gnou</i>	Black wildebeest	157	G	Short-grass grasslands	
<i>Elephas maximus</i>	Asian elephant	3270	G/B	Tropics	Ecological engineer in native range by dispersing large seeds and removing trees (Donlan et al. 2006)
<i>Equus asinus</i>	Donkey	180	G/B	Deserts	Digs wells used by other species
<i>Equus caballus</i>	Horse	400	G	Grasslands, open forests	Feeds on coarse, abrasive grasses (Naundrup & Svenning 2015)
<i>Hippopotamus amphibius</i>	Hippopotamus	1536	G	Aquatic daytime refuge; grasslands	Maintain grazing meadows, fertilize riparian systems (Bakker et al. 2016b), unstudied in introduced range
<i>Hippotragus niger</i>	Sable antelope	236	G/B	Woodland edges	
<i>Kobus ellipsiprymnus</i>	Waterbuck	204	G	Savanna woodlands	Riparian grazer, likely influences riparian vegetation and river geomorphology (IUCN 2017, Naiman & Rogers 1997)
<i>Oryx gazella</i>	Gemsbok	188	G	Desert scrub, desert grassland	Dig wells used by other species (Hamilton et al. 1977)
<i>Ovibos moschatus</i>	Muskox	313	G	Arctic tundra	Few other herbivores adapted to extreme arctic environment (Schmidt et al. 2015)
<i>Ovis ammon</i>	Argali	114	G	Steep, rocky environments	

<i>Rangifer tarandus</i>	Reindeer	109	G/B	Mountains, arctic tundra	Grazing can alter arctic albedo, causing temperature reductions that may counteract climate change (te Beest et al. 2016). Uniquely capable of digesting lichens (Palo 1993).
<i>Rucervus duvaucelii</i>	Barasingha	171	G	Forests, riparian grasslands	Riparian grazer, likely influences riparian vegetation and river geomorphology (IUCN 2017, Naiman & Rogers 1997)
<i>Rusa unicolor</i>	Sambar	178	G/B	Generalist	

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635 **FIGURE CAPTIONS**

636

637 Figure 1. Threatened megafauna species are finding refuge outside their native ranges.  
638 Percentage of megafauna in each family with introduced populations, colored by IUCN  
639 threat categories in their native ranges. Number within parentheses indicates total number  
640 of megafauna within each family.

641

642 Figure 2. The number of introduced megafauna species by IUCN (2017) threat status and  
643 population trends in their native ranges. The majority (59%) of introduced megafauna are  
644 threatened or have declining populations in their native ranges.

645

646 Figure 3. Percent of global populations of megafauna that are introduced. Color indicates  
647 IUCN (2017) status. Bars indicate high and low estimates if multiple estimates were  
648 found. Includes only species with known population sizes in native and non-native ranges  
649 and thus excludes *Tragelaphus eurycerus* (NT), *Kobus ellipsiprymnus* (LC), *Hippotragus*  
650 *niger* (LC), *Rusa unicolor* (VU), *Rucervus duvaucelii* (VU), *Alces alces* (LC), and *Ovis*  
651 *ammon* (NT). \* indicates post-domestic species.

652

653 Figure 4. Contribution of introduced megafauna to TDWG-country species richness. (a)  
654 Native megafauna species richness (b) introduced megafauna species richness, (c) all  
655 megafauna species richness, and (d) percent contribution of introduced species to  
656 TDWG-country megafauna assemblages. Inter- and intra-continental introductions were  
657 included. Native richness was derived from IUCN (2017) species distribution data.  
658 Introduced species distributions are available in Supplementary materials Appendix 1,  
659 Data A1.

660

661 Figure 5. Megafauna species richness per epoch by continent. 'Extinct' indicates species  
662 that went extinct in the wild on all continents; 'extirpated' are species that survived  
663 elsewhere; 'immigrated' are species that immigrated without human intervention;  
664 'introduced' indicates species introduced by humans; 'introduced, threatened' are  
665 introduced species threatened in their native ranges; 'survived' are species that were still  
666 present into the following epoch; 'survived, threatened' are threatened native species  
667 (Supplementary materials Appendix 3, Table A2).

668

669 Figure 6. Wild donkeys (*Equus asinus*) increase surface water availability in the Sonoran  
670 Desert. (a) Wild donkey digging well to water table ('burro well'), (b) troop of javelina  
671 (*Pecari tajacu*) bathing and drinking in burro wells, and (c) several-year-old Fremont's  
672 cottonwood (*Populus fremontii*) growing in an abandoned burro well on a high channel  
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681 SUPPLEMENTARY MATERIAL APPENDIX CAPTIONS

682 Supplementary material Appendix 1, Data A1. Shapefile of introduced megafauna ranges  
683 with references.

684

685 Supplementary material Appendix 2, Table A1. The introduced large ( $\geq 100\text{kg}$ )  
686 herbivorous mammals of the world. Data from Ripple *et al.* (2015), Long (2003), and the  
687 Large Herbivore Network (2015) unless otherwise noted.

688

689 Supplementary material Appendix 3, Table A2. Large ( $\geq 100\text{kg}$ ) herbivore species status  
690 by continent (Africa, Australia, Eurasia, North America, South America), by epoch  
691 (Pleistocene, Holocene, Anthropocene).

692

693 Supplementary material Appendix 4, Movie A1. Digging by wild donkeys ('burros')  
694 creates water resources used by other species in the Sonoran Desert. In addition to the  
695 four species shown here, from trail cameras and direct observations we have documented  
696 an additional twenty-six vertebrate species utilizing 'burro wells' including, bobcats  
697 (*Lynx rufus*), badgers (*Taxidea taxus*), black-tailed jackrabbits (*Lepus californicus*),  
698 striped skunks (*Mephitis mephitis*), ringtail cats (*Bassariscus astutus*), rock squirrels  
699 (*Spermophilus variegatus*), Common Ravens (*Corvus corax*), Hooded Orioles (*Icterus*  
700 *cucullatus*), Yellow Warblers (*Setophaga petechia*), Black-throated Sparrows  
701 (*Amphispiza bilineata*), Gila Woodpeckers (*Melanerpes uropygialis*), Gambel's Quail  
702 (*Callipepla gambelii*), Mourning Doves (*Zenaida macroura*), and various plant,  
703 amphibian and invertebrate species.