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Megaherbivores and cattle alter edge effects around ecosystem hotspots in an African savanna

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24 ABSTRACT

25 Wild mammalian herbivores and cattle are fundamental drivers of African savanna 26 ecosystems and have strong impacts on woody vegetation. However, few experimental 27 studies have investigated the separate and combined influences of different large herbivores 28 on spatial vegetation patterning. In East Africa, temporary cattle corrals (bomas) develop 29 after abandonment into productive, treeless 'glades' that attract both domestic and wild 30 herbivores. Edges of glades exhibit unusually high densities of large trees. We used a long-31 term, broad-scale manipulative experiment to test whether megaherbivores (elephants and 32 giraffes), wild meso-herbivores (15-1000 kg), or cattle caused shifts in woody plant 33 abundances in glade edges. We also examined cascading effects of megaherbivore and cattle 34 exclusion on symbiotic Acacia ants and wild meso-herbivores in glade edges. Megaherbivore 35 exclusion resulted in increased densities of tall trees, reproductive trees, and non-aggressive 36 Acacia ant species in glade edges. Cattle presence reduced meso-herbivore use inside and 37 away from glades, but not in glade edges. Our results suggest that megaherbivores and cattle 38 can dampen the magnitude of spatial patterns associated with glades and glade edges. These 39 findings provide insight into the development and maintenance of spatial heterogeneity in 40 savannas, and emphasize that land use change and mammalian extinctions have complex, 41 cascading ecological consequences.

42 Keywords: spatial heterogeneity; Acacia drepanolobium; Crematogaster species;

- 43 *Tetraponera penzigi*; pastoral; livestock-wildlife interactions
- 44

45 **1. INTRODUCTION**

In patchy and fragmented landscapes, ecological edge effects can have major impacts on
ecosystem structure and functioning, biotic interactions, and management (Fagan et al., 1999;
Ries et al., 2004; Harper et al., 2005). Edge effects are interactions between two adjacent

49 ecosystems or land cover types separated by an abrupt transition (Murcia, 1995). In savannas
50 as in forests, edge effects are often associated with changes in the composition and structure
51 of woody plant communities (Young et al., 1995; e.g., Brits et al., 2002; Lima-Ribeiro,
52 2008). In Brazilian cerrado fragments, for example, woody plants near fragment edges are
53 shorter and thinner than plants in fragment interiors (Lima-Ribeiro, 2008). Edge-related
54 changes in woody plant structure and composition can have cascading effects on other
55 species and processes (Fagan et al., 1999; Porensky, 2011).

56 Large mammalian herbivores shape African savanna ecosystems and have strong impacts 57 on woody vegetation (e.g., Pellew, 1983; Augustine and McNaughton, 2004; Goheen et al., 58 2010; Porensky and Veblen, 2012). Thus, in these landscapes, large herbivores are likely to 59 influence edge effect patterns associated with woody vegetation. However, few experimental 60 studies have examined whether edge effects are altered by large herbivores (but see Fox et 61 al., 1997; Allombert et al., 2005; Didham et al., 2009), and to our knowledge, none has done 62 so in Africa where large herbivores are still abundant and diverse. Further, to our knowledge, 63 the separate and combined impacts of different guilds of large herbivores on edge effects 64 have not been experimentally tested (but see observational work by Brits et al., 2002; de Beer 65 et al., 2006). With a better understanding of how large herbivores alter edge effects, 66 ecologists will be able to better comprehend the maintenance of spatial heterogeneity in 67 savannas, as well as the potential landscape-level consequences of mammalian extinctions. 68 We used a long-term, broad-scale manipulative experiment to determine how various 69 large herbivores alter edge effects around anthropogenic nutrient hotspots. For centuries, 70 pastoralists in sub-Saharan Africa have corralled their cattle in temporary thorn-fence 71 'bomas' at night for protection against predators and stock raiders (Western and Dunne, 72 1979; Blackmore et al., 1990). Bomas are typically 50-100 m in diameter and are used for 73 months or years before being abandoned. After abandonment, dung-filled boma sites develop

into distinctive ecosystem 'hotspots' that persist for decades to centuries (Blackmore et al.,
1990; Young et al., 1995; Augustine, 2003). At our study site, abandoned bomas develop into
treeless, productive 'glades' that harbor nutrient-rich plant species and are used preferentially
by livestock and wild herbivores (Young et al., 1995; Augustine, 2003, 2004; Veblen and
Young, 2010; Porensky, 2011; Veblen, 2012). Glades are common in this landscape (found at
densities of about 2 glades per km², Veblen, 2012), and represent major sources of structural
and functional heterogeneity.

81 Glades can have edge effects that extend at least 100 m into the surrounding savanna 82 matrix (Young et al., 1995; Porensky, 2011). One striking pattern associated with glade edges 83 is an unusually high density of large trees (Porensky, 2011), which might be related to high 84 nutrient availability within glades (Veblen, 2012) or a competitive advantage gained by tree 85 seedlings growing in heavily grazed areas (Riginos and Young, 2007). Large trees around 86 glades might also reflect the legacy of an establishment phase that occurred during or soon 87 after active boma use (Muchiru et al., 2009). In this study, we investigated impacts of 88 different large herbivores on woody vegetation patterns in glade edges. Large herbivores are 89 known to affect the long-term development of herbaceous communities in glades (Veblen, 90 2008; Veblen and Young, 2010), but their impacts on co-occurring woody plants remain 91 unclear (but see Porensky and Veblen, 2012).

Ecological edges can have strong influences on the interactions among different associated species (Fagan et al., 1999). Our study system gave us the opportunity to experimentally investigate how large herbivores, by altering vegetation in glade edges, may have cascading impacts on other animal taxa. For example, by thinning tall trees around glades, elephants may decrease the use of glade edges by other herbivores that respond to the micro-climatic benefits and high-quality forage found beneath trees (Ludwig et al., 2008; Treydte et al., 2009). Conversely, reduced tree densities may afford increased visibility and

99	enhanced predator detection, which might be attractive to mid-sized herbivores (Riginos and						
100	Grace, 2008). Further, large herbivores could affect invertebrate communities associated with						
101	glade edges. In particular, the presence or absence of megaherbivore browsers might alter the						
102	relative abundances of ant species that are symbiotic with the dominant Acacia tree species in						
103	our system (Young et al., 1997).						
104	This study used a broad-scale manipulative experiment to test three hypotheses about large						
105	herbivores and glade edge effects.						
106	1) Woody vegetation patterns in glade edges are significantly altered by large						
107	herbivores.						
108	2) Different guilds of large herbivores (e.g., cattle vs. megaherbivores) have different						
109	impacts on woody vegetation patterns.						
110	3) Impacts of large herbivores on woody vegetation have cascading consequences for						
111	mid-sized herbivores and symbiotic Acacia ants.						
112	2. METHODS						
113	2.1 Study area						
114	This study took place at the Mpala Research Centre in Laikipia County, Kenya (0°17`N,						
115	36°52'E, 1800 m asl), where more than 2,000 Boran cattle (Bos indicus) coexist with						
116	abundant wildlife on a 17,000 ha conservancy. The climate is semi-arid with a mean annual						
117	rainfall of 500-600 mm (Young et al., 1998).						
118	The study site is located on "black cotton" vertisol soil that supports some of the most						
119	productive rangelands in Africa (Young et al., 1998). Large mammalian herbivores are						
120	abundant; species include the endangered Grevy's zebra (Equus grevyi), the more common						
121	Burchell's zebra (Equus quagga burchelli), eland (Taurotragus oryx), oryx (Oryx beisa),						
122	hartebeest (Alcelaphus buselaphus), steinbuck (Raphicerus campestris), Grant's gazelle						
123	(Nanger [Gazella] granti), warthog (Phacochoerus africanus), and occasionally bush duiker						

124 (Sylvicapra grimmia) or impala (Aepyceros melampus). These species are categorized as 125 meso-herbivores hereafter. The study site also supports low densities of African buffalo 126 (Syncerus caffer), which are not included in meso-herbivore analyses since their dung piles 127 are indistinguishable from those of cattle. In addition, two megaherbivore species, giraffe 128 (Giraffa camelopardis reticulata) and African elephant (Loxodonta africana), are present. 129 The dominant woody species in this system is Acacia drepanolobium. This small tree 130 grows up to 6 m tall and accounts for 97% of total woody cover at our study site (Young et 131 al., 1998; Riginos and Grace, 2008). Individuals produce hollow swollen spines that are 132 inhabited by symbiotic ants, which defend the tree against herbivory (Young et al., 1997; 133 Young and Okello, 1998; Palmer et al., 2010).

134 In our study system, A. drepanolobium trees are typically inhabited by one of four species of ants: Tetraponera penzigi, Crematogaster mimosae, C. nigriceps, or C. sjostedti (Young et 135 136 al., 1997; Stanton et al., 2002). These ants exist within a competitive hierarchy associated 137 with differently-sized trees. Tetraponera penzigi and C. nigriceps are competitively 138 subordinate and tend to be found on smaller trees, while C. mimosae and C. sjostedti are 139 competitively dominant and often found on larger trees (Young et al., 1997; Stanton et al., 140 2002; Palmer et al., 2010). The four species also differ in their ability to defend trees against 141 megaherbivore browsing: Tetraponera penzigi and C. sjostedti ants are less aggressive, while 142 C. mimosae and C. nigriceps are more aggressive towards browsers (Young et al., 1997; 143 Stanton et al., 2002; Young et al., 2003; Palmer et al., 2010). Hereafter, we refer to ants based 144 on these two aggressiveness classes.

Our work took place within the Kenya Long-term Exclosure Experiment (KLEE), which has been assessing the separate and combined impacts of different herbivores on this savanna ecosystem since 1995 using an exceptionally broad-scale approach (Young et al., 1998). The experiment consists of three replicate blocks, each with six 200 x 200 m treatment plots (Fig. 149 1). Different barriers exclude different guilds of herbivores (see also Appendix): 1) Visual 150 markers serve to keep individually-herded cattle (C) in designated plots. 2) A two-strand 151 electric line 2 m off the ground excludes only megaherbivores (elephants and giraffes; M). 3) 152 A nine-strand electric game fence excludes all meso-herbivores 15-1000kg (W) as well as 153 megaherbivores. There are gates in the game fence to allow entry by herded cattle. In 154 addition to total exclusion plots (O), treatment combinations include C, W, WC, MW and 155 MWC, where letters indicate the types of animals allowed into the plots (e.g., WC plots allow 156 wildlife and cattle). Smaller herbivores (steinbuck, hares, rodents, and invertebrates) have 157 access to all plots. Individually-herded groups of cattle were run in each C, WC or MWC plot 158 for 2 hours 6-8 times per year, producing impacts similar to the overall ranch stocking density 159 (see Odadi et al., 2007) while limiting human and livestock presence to a few hours per year. 160 The KLEE study design includes six glades (abandoned boma sites), which were divided 161 in 1995 by the herbivore exclusion fences to create 18 glade portions (Fig. 1a, Young et al., 162 1998). These include several glade portions adjacent to the KLEE plots that are maintained in 163 herbivore treatments (Fig. 1a, Young et al., 1998) which results in comparable environmental 164 conditions within each portion. Historical aerial photographs indicate that all of these glades 165 have been present for more than 40 years.

166 2.2 Data collection

Over 2006-2007, we determined whether woody communities in glade edges differed from background conditions across herbivore treatments. For all glade portions in KLEE (Fig. 1a), we sampled woody species in large plots at the glade edge and at 150 m away from the glade. At 150 m outside the perimeter of each of the 18 glade portions, we placed the closest, long edge of a 40 x 30 m plot (Fig. 1b). At glade edge, we split sampling into two 20 x 30 m plots to increase the amount of curvilinear glade perimeter captured by our plots. Each 20 x 30 m plot was placed at glade edge, spanning 10 m inside to 20 m outside the glade perimeter (Fig. 174 1b). We counted all *A. drepanolobium* trees rooted in each sample plot and categorized them
175 into three height classes: <1 m, 1-3 m, and >3 m tall. We also recorded all occurrences of
176 non-*A. drepanolobium* woody species (excluding seedlings).

177 In 2009 (a drought year), we looked more closely at A. drepanolobium, collected edge 178 data with higher spatial resolution, and investigated meso-herbivore use of glades and glade 179 edges. The 2006-2007 data suggested that meso-herbivores had no consistent effects on 180 woody vegetation patterns around glades (see Results). Therefore, in 2009 we concentrated 181 on the effects of cattle and megaherbivores. We used four of the six KLEE treatments (W, 182 WC, MW, and MWC) to investigate a full factorial of megaherbivore and cattle effects. The 183 2006-2007 data also revealed a strong pattern in the largest size class of A. drepanolobium 184 trees (>3 m). In 2009, we further subdivided this large size class into 3-4 m and >4 m size 185 classes to improve the resolution of our data. For all size classes we also included metrics of 186 A. drepanolobium reproduction and symbiotic ant occupancy. Finally, we used dung counts 187 to investigate interactions within the herbivore community.

For all 2009 surveys, we set up 150 m-long transects at each glade inside or adjacent to KLEE treatment plots. Each transect started in the center of a glade and extended away from the glade and into one of the treatment plots (Fig. 1a). Along each transect, we recorded the following response variables:

(1) For each *A. drepanolobium* tree within 4 m of the transect line, we recorded its
distance along the transect, its height, and its reproductive status (flowering or fruiting
vs. non-reproductive). Trees were classified into four height categories: <1 m, 1-3 m,
3-4 m, and >4 m. We also recorded the species identity of any symbiotic ant species
present on each tree.

197 (2) For each dung pile within 4 m of the transect line, we recorded its distance along the
198 transect and the animal species that produced it. Previous work suggests that dung

counts are a reliable method to determine relative animal use (Augustine 2003, Young
et al. 1995, 2005). Buffalo and cattle dung piles were indistinguishable, and therefore
these two species were lumped as 'cattle+buffalo'. However, cattle are >10 times
more abundant than buffalo at the study site. Other wildlife were classified as
'grazers' (zebra, hartebeest and warthog), 'mixed feeders/browsers' (eland, oryx,
steinbuck, Grant's gazelle, bush duiker and impala), or 'megaherbivores' (elephant
and giraffe).

For each of these response variables, data were binned into 5 m distance intervals along eachtransect.

208 2.3 Statistical analyses

209 2.3.1 2006-2007 surveys

210 We used split-plot designs to test treatment effects on woody vegetation. For woody (non-A. 211 drepanolobium) species densities and the three A. drepanolobium size class densities, each 212 model included the following fixed effects: main plot effect of herbivore treatment (O, C, W, 213 WC, MW, MWC), subplot effect of distance (inside, edge, outside of glade), and the 214 herbivore*distance interaction. Block, glade nested within block, and the block*glade 215 interaction were included as random effects. For all statistical analyses, we used generalized 216 linear mixed models (GLMMs), maximum-likelihood methodology and Satterthwaite's 217 approximation of degrees of freedom (PROC MIXED, version 9.1, SAS Institute 2002). 218 Variance-weighting was used when variances were not homogenous, and values were log-219 transformed when necessary. We used Tukey's HSD analyses for post-hoc comparisons, and 220 we report means ± 1 SE throughout the results. 221 2.3.2 2009 surveys: standardizing the glade edge

222 One of the most defining structural characteristics of glades is their persistent lack of trees, so

tree density was used to standardize the location of the glade edge across transects. Although

this 'glade edge' location is not necessarily intended to reflect a functional edge, our observations suggest it is largely coincident with the location of the original boma fence. The glade edge was defined as the first 5 m interval in which we found $\geq 4 A$. *drepanolobium* trees/40 m². For each transect, this glade edge distance was defined at 0 m, so that areas inside and outside the glade had negative and positive distance values, respectively.

229 2.3.3 2009 surveys: spatially coarse-scale analysis

230 For each response variable along each transect, we averaged binned values for two 231 standardized distance intervals: -25 to 0 m (inside glade), and 0 to 25 m (glade edge). For 232 each distance interval, we compared among treatments using GLMMs with the following 233 predictors and their interactions: block (a random factor), cattle exclusion, and 234 megaherbivore exclusion. The cattle*megaherbivore interaction term was removed from 235 models in which it was non-significant (p > 0.10). Response variables included overall tree 236 density, tree density separated by size class, overall density of meso-herbivore dung piles, 237 meso-herbivore dung density separated by feeding guild, overall density of trees occupied by 238 Acacia ants, and ant occupancy separated by species aggressiveness. For analyses involving 239 multiple herbivore guilds, tree size classes or ant species, MANOVAs were used initially to 240 establish overall significance. For all other analyses, we used GLMMs with maximum-241 likelihood methodology and Satterthwaite's approximation of degrees of freedom (PROC 242 MIXED, version 9.1, SAS Institute 2002). Response variables were log-transformed when necessary to achieve normality. Due to the low replication of this broad-scale experiment, we 243 244 set α at 0.10. We also used a generalized linear mixed model (with predictors as described 245 above and log-transformed glade radius as the response variable) to confirm that cattle and megaherbivore exclusion had no significant impacts on glade radius. We report means ± 1 SE 246 247 throughout the results.

248 2.3.4 2009 surveys: spatially fine-scale analysis

For response variables significantly affected by cattle or megaherbivore exclusion (based on the coarse-scale analysis above), we fit continuous nonlinear models to identify the distance ranges over which significant differences occurred. We used the following model, which includes linear, sigmoid, and unimodal edge effect shape components:

253
$$y=a+b \times \frac{b}{1+e^{(b_2-x)b_3}} + h^* e^{\frac{-(x-x_0)^2}{2W^2}}$$

254 linear sigmoid unimodal

255 where *X* is distance from the edge and the other variables are fitted constants. For each 256 response variable along each transect, we fitted the model using the nonlinear platform in 257 JMP (version 8.0, SAS Institute, Inc., Cary, NC, USA) and an expectation maximization 258 approach (for methodological details, see Porensky, 2011). We averaged fitted models within 259 different treatments and then identified regions in which the 90% confidence intervals of the 260 different average models were non-overlapping. In these regions, treatments were considered 261 significantly different. To reduce the influence of outliers, we only compared standardized 262 distance values at which at least four transects from each treatment had data. The model-263 fitting and subsequent model-averaging process should further reduce the influence of outliers. 264

265 **3. RESULTS**

266 3.1 2006-2007 surveys

267 Densities of the largest individuals (>3 m tall) of the dominant tree, Acacia drepanolobium,

268 were three times higher in glade edges than far from glades (Table 1, distance effect $F_{1,12,2}$ =

269 57.34, *p* < 0.0001). However, this effect occurred only in plots without megaherbivores

270 (edge>out, treatment*distance class interaction $F_{5,12,3}$ = 3.42, p= 0.04, Tukey HSD p <0.05 for

271 W, WC, and C; for O p=0.19). In MW and MWC plots, the two plot types that allow entry to

272 megaherbivores, edges did not have significantly higher densities (Table 1). For 1-3 m tall A.

273 drepanolobium trees there were no significant distance or treatment effects (Table 1). For 274 small (<1 m tall) trees, no significant distance effects could be found. The C plots (cattle 275 only) had 39-60% fewer small A. drepanolobium individuals than the other treatments, but 276 this was statistically significant only compared to WC (meso-herbivores + cattle, Table 1, 277 treatment effect $F_{5, 15.8}$ = 2.92, p= 0.05). In other words, plots allowing only cattle had fewer 278 small trees than plots allowing both meso-herbivores and cattle. However, small tree density 279 did not differ significantly between total exclusion plots (O) and plots allowing meso-280 herbivores (W).

281 Densities of two common woody shrubs, *Lycium europaeum* L. and *Rhus natalensis* 282 Krauss, were significantly higher in glade edges than outside glades (*L. europaeum*: 44.1 ± 283 13.9 ha⁻¹ vs. 3.9 ± 3.0 , $F_{1,19,1} = 5.83$, p = 0.03; *R. natalensis*: 16.2 ± 4.1 vs. 5.4 ± 1.2 , $F_{1,14} =$ 284 7.95, p = 0.01). Densities of the other most common shrub, *Cadaba farinosa* Forssk., did not 285 differ significantly by distance (27.9 ± 4.1 vs. 20.1 ± 5.4 , $F_{1,11,2} = 1.98$, p = 0.19). None of 286 these shrubs responded significantly to herbivore treatments.

287 3.2 2009: Transect-wide meso-herbivore results

Averaged across treatments (W, WC, MW and MWC) and distances from glades, mesoherbivore (excluding cattle and buffalo) dung comprised 63% zebra, 19% eland, 6% oryx, 5% hartebeest, 4% steinbuck, and 3% other species. In plots where cattle were allowed, cattle+ buffalo dung and zebra dung each accounted for $37 \pm 6\%$ of total dung. In plots where cattle were excluded, cattle+buffalo dung accounted for $3 \pm 1\%$ and zebra dung accounted for $60 \pm$ 4% of total dung. Zebra dung was 70% more abundant in plots where cattle were excluded (129 ± 16 vs. 222 ± 16 dung piles; t = -4.11, p = 0.003, df = 8.8).

295 3.3 2009: Coarse-scale analysis

296 Cattle and megaherbivore exclusion had no significant impacts on glade radius (W= 35 ± 9

297 m, WC= 28 ± 3 m, MW= 28 ± 3 m, MWC= 27 ± 4 m; all *p*-values >0.10). In glade edges,

298 herbivore treatments significantly affected tree size structure (tree size class MANOVA: 299 cattle effect Wilk's $\lambda = 0.04$, p = 0.02; megaherbivore effect Wilk's $\lambda = 0.05$, p = 0.03). In 300 particular, the density of large trees (>4 m tall) in glade edges was more than three times higher when megaherbivores were excluded (Fig. 2; 2.0 ± 0.2 vs. 0.6 ± 0.2 trees/ 40 m²; 301 302 megaherbivore $F_{1.5,29}$ = 50.44, p= 0.0007). Cattle exclusion did not significantly affect large 303 tree density in glade edges ($F_{1,3,74}$ = 1.43, p= 0.3). In glade edges, neither cattle nor 304 megaherbivore exclusion significantly affected the density of trees in other size classes (Fig. 305 2) or overall tree density (all *p*-values >0.10).

306 Cattle exclusion and megaherbivore exclusion both led to higher densities of

307 reproductively active trees in glade edges. When cattle were excluded, the density of 308 reproductively active trees was 80% higher $(3.2 \pm 0.6 \text{ vs}, 1.8 \pm 0.3 \text{ flowering or fruiting trees})$ 40 m²; $F_{1.8}$ = 9.42, p= 0.02). Similarly, the density of reproductively active trees was over 309 310 twice as high when megaherbivores were excluded (Fig. 2; 3.2 ± 0.6 vs. 1.5 ± 0.2 flowering or fruiting trees/ 40 m²; $F_{1.8}$ = 13.25, p= 0.007). 311

312 Cattle and megaherbivore exclusion affected the densities of trees occupied by different 313

types of ants in glade edges (cattle*megaherbivore effect *Wilk's* $\lambda = 0.27$, p = 0.07).

314 Megaherbivore exclusion resulted in higher densities of non-aggressive ants (T. penzigi and

315 C. sjostedti), especially when cattle were also excluded (Fig. 3; cattle $F_{1.7}$ = 4.93, p= 0.06;

316 megaherbivore $F_{1,7}$ = 11.89, p= 0.01; cattle*megaherbivore $F_{1,7}$ = 4.43, p= 0.07). Neither cattle

317 nor megaherbivore exclusion significantly affected the density of trees occupied by more

318 aggressive ants (C. nigriceps and C. mimosae; p-values >0.10). Similarly, the overall density

319 of ant-occupied trees in glade edges was not significantly affected by either cattle or

320 megaherbivore exclusion (*p*-values >0.10).

321 Inside glades, cattle exclusion significantly affected the density of wildlife dung piles

322 (wildlife MANOVA: cattle effect *Wilk's* $\lambda = 0.32$, p = 0.06; megaherbivore effect *Wilk's* $\lambda =$ 323 0.64, p=0.3). For grazers (zebra, hartebeest and warthog), the density of dung piles inside 324 glades was more than twice as high when cattle were excluded (10.8 ± 1.3 vs. 4.7 ± 1.0 325 grazer dung piles per 40 m²; cattle effect $F_{1,8}=19.45$, p=0.002). Dung densities for browsers 326 and mixed feeders (eland, oryx, steinbuck, Grant's gazelle, bush duiker and impala) inside 327 glades were also more than twice as high when cattle were excluded (3.8 ± 0.6 vs. 1.7 ± 0.5 328 browser/mixed feeder dung piles per 40 m²; cattle effect $F_{1,8}=6.45$, p=0.03).

329 Megaherbivore exclusion did not significantly affect the density of wildlife dung piles 330 inside glades (grazers: $F_{1,8}$ = 2.97, p= 0.12; browsers/mixed feeders: $F_{1,8}$ = 0.08, p= 0.8).

331 Moreover, neither cattle nor megaherbivore exclusion significantly impacted the density of

332 wildlife dung piles in glade edges (MANOVA *p*-values >0.10). Inside glades, neither cattle

333 nor megaherbivore exclusion significantly affected overall tree density, the density of

reproductively active trees, or the density of ant-occupied trees (all *p*-values >0.10).

335 Separating trees by size class and separating ants by aggressiveness did not reveal any

additional patterns inside glades (MANOVA *p*-values >0.10).

337 3.4 2009: Fine-scale analysis

338 To examine herbivore effects at a finer spatial scale, we compared average fitted models 339 obtained from transects in different treatments. Average models were considered significantly 340 different wherever their 90% confidence intervals did not overlap. For grazer use, browser 341 use, reproductively active trees and trees occupied by non-aggressive ants, we compared 342 fitted models from transects in plots with or without cattle. At distances <-20 m, between -20 343 and 0 m, and between 50 and 90 m outside glade edge, grazer use was significantly higher in 344 transects without cattle (Fig. 4a). At distances between -15 and 5 m, browser use was also 345 significantly higher in transects without cattle (Fig. 4b). Transects without cattle had 346 significantly more reproductively active trees between 5 and 15 m outside glade edge, and 347 significantly more non-aggressive ants between 40 and 50 m outside glade edge.

348 For densities of large trees (>4 m tall), reproductively active trees and trees occupied by 349 non-aggressive ants, we compared fitted models from transects in plots with or without 350 megaherbivores. When megaherbivores were excluded, large tree densities were significantly 351 higher at distances between -10 and 10 m, 15-35 m, and 65-80 m outside glade edge (Fig. 352 4c). Transects without megaherbivores also had significantly more reproductively active trees 353 between 20 and 35 m (Fig. 4d). These results indicate that megaherbivore exclusion led wider 354 peaks for large and reproductively active tree densities in glade edges (Fig. 4c,d). 355 Megaherbivore exclusion did not significantly affect non-aggressive ant occupancy at any 356 distance, likely because of the interaction between cattle and megaherbivore effects (see 357 above, Figs. 3 and 4e).

358 4. DISCUSSION

359 Megaherbivores and cattle both significantly altered glade edge effects in this savanna 360 landscape mosaic. Megaherbivores reduced the spatial heterogeneity created by glades, not 361 by reducing differences between glades and the background savanna, but by reducing unique 362 traits associated with glade edges: high densities of large trees, reproductively active trees 363 and non-aggressive Acacia ants. Cattle also reduced the densities of reproductively active 364 trees and non-aggressive Acacia ants in glade edges. Cattle further dampened spatial 365 heterogeneity by weakening meso-herbivore preference for glade interiors. These results 366 reflect a combination of direct and indirect effects (Fig. 5), and further research is needed to 367 clarify some of the relevant mechanisms.

Meso-herbivore exclusion did not have strong impacts on *A. drepanolobium* density or size structure in glade edges. Cattle probably have stronger impacts than wild mesoherbivores because of their higher biomass density (cattle have 5–10 times higher biomass per km² than wild ungulates). Differences between meso-herbivores and cattle might also be related to differences in the timing of grazing (continuous vs. discontinuous), evolutionary

history in the system (long vs. short), or diet (though dietary overlap between cattle and the
dominant meso-herbivore, zebra, is probably large) (Odadi et al., 2007; Odadi et al., 2011).

375 4.1 Acacia drepanolobium

376 One of the most striking edge effects in our study system is the high density of large trees in 377 glade edges. Although it is not clear what causes rings of tall, dense trees to form at glade 378 edges, we found that megaherbivore exclusion clearly enhanced this effect. The abundance of 379 large trees in glade edges was significantly higher in plots where megaherbivores were 380 excluded (Table 1; Figs. 2, 4 and 5). This suggests that megaherbivores preferentially feed on 381 large trees in glade edges, perhaps because higher tree densities and taller trees in glade edges 382 represent attractive forage patches. Elephants often seem to browse preferentially on larger 383 trees, especially in high tree density areas (Pellew, 1983; Augustine and McNaughton, 2004). 384 Trees next to glades may also contain more nutrients than trees in the background savanna, 385 and this enhanced nutrient content may attract elephant and giraffe browsing. Elephant 386 foraging typically leads to killing, coppicing or reduction of tree size (Van de Vijver et al., 387 1999; Augustine and McNaughton, 2004; Goheen et al., 2007) while giraffe browsing tends 388 to be focused on taller trees and can severely slow tree growth (Pellew, 1983).

389 Whereas only megaherbivores had a negative effect on tree density in glade edges, both 390 megaherbivores and cattle reduced fruiting and flowering of A. drepanolobium (Fig. 5). 391 These results may be driven by changes in browsing pressure. Megaherbivore exclusion 392 reduces browsing directly, and cattle exclusion may also reduce browsing pressure by 393 increasing understory forb biomass (Odadi et al., 2013). In cattle exclusion plots, wild 394 ungulates may able to meet nutritional needs by eating forbs, and therefore may be less likely 395 to browse on trees. Reduced browsing can increase reproduction by increasing tree size, since 396 larger trees are more likely to reproduce (Goheen et al., 2007; Palmer et al., 2010). Reduced 397 browsing can also cause reduced investment in defense (Young et al., 2003; Palmer et al.,

2008), which may lead to increased investment in reproduction (Goheen et al., 2007). Finally,
increased reproduction in the absence of megaherbivores and cattle could be an indirect
consequence of increased densities of non-aggressive ants. *Crematogaster sjostedti*occupation is associated with increased fruiting in *A. drepanolobium*, and *T. penzigi* is also
associated with higher fruiting probabilities than either *C. mimosae* or *C. nigriceps* (Palmer et
al., 2010).

404 If increased flowering and fruiting near glades leads to higher seedling recruitment inside 405 glades, herbivore exclusion might reduce the persistence of glades as treeless areas (by 406 promoting tree reproduction). Alternatively, higher seed availability in the absence of large 407 herbivores might enhance granivorous insect and rodent populations, which in turn could 408 suppress seedling establishment and keep glades treeless (Walters et al., 2005; Palmer and 409 Brody, 2007). Moreover, relationships between the presence of reproductive structures and 410 actual seed production or seedling recruitment are probably highly variable (Goheen et al., 411 2007; Goheen et al., 2010). In light of these complications, impacts of different herbivores on 412 tree recruitment inside and near glades warrant further study.

413 4.2 Symbiotic Acacia ants

Our study showed that the number of trees occupied by non-aggressive ants was higher when both megaherbivores and cattle were absent (Figs. 3 and 5). The dominant non-aggressive ant species (*C. sjostedti*) is more common in larger trees (Young et al., 1997), so increased densities of large trees certainly contributed to increased densities of non-aggressive ants. However, changes in large tree density are not sufficient to explain changes in ant occupancy (Fig. 4), and we found that megaherbivore and cattle exclusion did not significantly affect small- or medium-sized tree density in glade edges.

421 Megaherbivore and cattle exclusion may further affect changes in ant occupancy by
422 reducing browsing pressure (see section 4.1). Trees under lower herbivory pressure can

reduce production of 'ant rewards' in the form of nectaries and swollen spines (Huntzinger et al., 2004; Palmer et al., 2008), causing ant species more dependent on rewards (e.g., *C. mimosae*) to be replaced by ant species less dependent on rewards (e.g., *C. sjostedti*; Palmer et al., 2008). Our results are similar to those of Palmer et al. (2008), who observed a
replacement of *C. mimosae* (aggressive) by *C. sjostedti* (non-aggressive) under herbivore exclusion.

429 Changes in ant community composition can have major impacts on *A. drepanolobium*430 survival, growth, architecture, parasitism and reproduction (Stanton et al., 1999; Palmer et al.,
431 2002; Riginos and Young, 2007; Palmer et al., 2008; Palmer et al., 2010). Our study shows
432 that these changes could have spatially explicit ecosystem effects, particularly in areas where
433 *A. drepanolobium* is dominant (Young et al. 1998).

434 **4.3** Mammal use

435 Cattle and zebra were the species whose dung was most frequently found at our study site; 436 these two species dominate this savanna landscape (Young et al., 2005). The 70% increase in 437 zebra dung in plots excluding cattle is consistent with previous studies demonstrating that 438 cattle and zebra, both grazers, usually compete for their diet and prefer similar habitat, 439 particularly during the dry season (Young et al., 2005; Odadi et al., 2007; Odadi et al., 2011). 440 Our data, collected during a drought, suggested that cattle reduced meso-herbivore 441 preference for glades (Fig. 4). Given that cattle spend only 12-16 hours per year inside each 442 glade portion, our results probably were not driven by direct interference between cattle and 443 wildlife. Though cattle had no significant effects on tree densities or size structure (Table 1), 444 cattle were associated with fewer reproductive trees and higher aggressive ant occupancy 445 (Fig. 3). By increasing aggressive ant occupancy, cattle could have made A. drepanolobium 446 trees less attractive to meso-herbivore browsers. However, our data suggest that cattle had 447 similar effects on meso-herbivore grazers and browsers. Thus, we hypothesize that the effects of cattle on wildlife were driven mostly by indirect competition for herbaceous forage (Fig. 5;Odadi et al., 2011).

450 In the presence of cattle, meso-herbivore dung levels were constant across the landscape, 451 suggesting a weak response of wildlife to glades. However, in the absence of cattle, meso-452 herbivore preference for glade interiors was pronounced. Preference of wildlife for glade 453 interiors has also been found in other landscapes in eastern Africa (Young et al., 1995; 454 Augustine, 2004; Treydte et al., 2006b) and has been attributed to factors such as high forage 455 quality (Augustine, 2003; Treydte et al., 2006a) or better visibility within glades (Young et 456 al., 1995; Riginos and Grace, 2008). In the absence of cattle, high meso-herbivore dung 457 densities inside glades were offset by relatively low dung densities in glade edges. This edge 458 pattern may reflect foraging decisions made by meso-herbivores approaching glades which, 459 in the absence of cattle, are full of palatable forage.

Previous research has shown that grazing by wild meso-herbivores helps maintain glades in a successional stage dominated by a highly palatable short-grass (Veblen and Young, 2010). This in turn further attracts wild herbivore grazing and may help maintain high glade nutrient levels over the long-term by offsetting nutrient export via herbivory with nutrient import via dung deposition. When cattle are present, these feedbacks may be weakened.

465 4.4 Conclusions

Our findings demonstrate that large mammalian herbivores can have profound impacts on landscape heterogeneity in an African savanna. Cattle and megaherbivores dampened the ecosystem heterogeneity created by treeless glades embedded within a savanna matrix. Although glade edges still retain high tree densities in the presence of megaherbivores, this pattern would be even more striking if megaherbivores were excluded. Similarly, our results suggest that wildlife preference for glade interiors would be even stronger in the absence of

472 cattle. Because glades are common features in livestock-dominated savanna landscapes, and

are now being actively managed for their ecosystem benefits (Porensky, Veblen, Riginos,
pers. observ.), our results can inform management decisions in areas where wildlife and
livestock share resources. More broadly, our results demonstrate that edge effects are
sensitive to the presence and activities of various types of large mammalian herbivores.
Future research could explore the nature and importance of this context-dependence in other
study systems, as well as its implications for biodiversity conservation in fragmented
landscapes.

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604

606 **7. TABLES**

607 **Table 1:** Mean (±1 SE, in trees per hectare) densities of three size classes of *Acacia*

608 *drepanolobium* trees in glade edges and 150 m outside of glades in six herbivore treatments.

- 609 Herbivore treatments allow different combinations of the following: cattle ("C"), wild meso-
- 610 herbivores >15 kg ("W"), and wild megaherbivores (elephants and giraffes, "M"). No
- 611 herbivores are allowed in "O" plots. Letters in the >3 m tree category indicate significant
- 612 differences across all 12 means (Tukey's HSD. α = 0.05). Differences were not significant in

613 the <1 m and 1-3 m categories.

T	Herbivore treatment	edge		150 m away	
Tree size class		mean (#/ha)	\pm SE	mean (#/ha)	± SE
	C	128	53	314	23
	MW	304	21	521	229
ш	MWC	461	69	264	84
$\overline{\lor}$	0	367	121	389	161
	W	286	66	725	168
	WC	442	17	686	225
	С	797	57	917	128
	MW	742	217	1433	608
ш	MWC	875	230	519	136
1-3	0	1339	167	1136	363
	W	1003	61	1253	287
	WC	1222	242	1075	393
	С	^a 653	129	^b 286	85
	MW	^{ab} 329	71	^b 208	50
Ш	MWC	^b 189	26	^b 131	31
>3	0	^{ab} 486	92	^b 225	97
	W	^a 864	220	^b 158	17
	WC	^a 664	266	^b 131	64

616 8. FIGURE LEGENDS

617 Fig. 1: a) Diagram of the Kenva Long-term Exclosure Experiment, including herbivore 618 treatment labels, locations of glade portions and 2009 sampling transect locations. See Table 619 1 for explanations of treatment abbreviations. Glade portions occurring in treatment plots 620 labeled in bold were sampled in 2006-2007, and b) provides a detailed view of 2006-2007 621 sample plot locations. 622 **Fig. 2:** Mean density (± 1 SE; in trees per 40 m²) of *A. drepanolobium* trees of different sizes 623 and reproductive states in glade edges (0-25 m distance class), in plots to which 624 megaherbivores did or did not have access. Asterisks indicate significant differences between 625 herbivore treatments at the p < 0.05 level. **Fig. 3**: Mean densities $(\pm 1 \text{ SE}, \text{ in trees per } 40 \text{ m}^2)$ of trees hosting non-aggressive ant species 626 627 (C. sjostedti and T. penzigi) in different large herbivore treatments. + M =plots allowing 628 megaherbivores; - M = plots excluding megaherbivores; + C = plots allowing cattle; - C = 629 plots excluding cattle. The interaction between cattle and megaherbivore presence was 630 significant at the p <0.10 level (cattle $F_{1,7}$ = 4.93, p= 0.06; megaherbivore $F_{1,7}$ = 11.89, p= 631 0.01; cattle*megaherbivore $F_{1.7}$ = 4.43, p= 0.07). 632 **Fig. 4:** Average fitted models (± 1 SE) for densities of a) grazer dung piles, b) browser dung 633 piles, c) large trees (>4 m tall), d) reproductively active trees, and e) non-aggressive ants. 634 Fig. 5: Conceptual diagram illustrating direct and indirect effects of herbivore treatments on 635 meso-herbivore use and A. drepanolobium density, reproductive status and ant occupancy. 636 White boxes indicate herbivore treatments and gray boxes indicate results from this study. 637 Solid arrows and black ovals indicate mechanisms supported by published research. Dotted 638 arrows and gray ovals indicate currently untested mechanisms. 639









Figure 2









652 **Figure 5**



Citations:¹Augustine and McNaughton, 2004;²Goheen et al., 2007;³Palmer et al., 2008; ⁴Young et al., 1997;⁵Palmer et al., 2010;⁶Odadi et al., 2011

654 Appendix: Abundance of cattle and megaherbivore dung

- 655 Dung measurements indicated that cattle and megaherbivore exclusion treatments were
- 656 effective. Inside glades, cattle + buffalo dung density was 10 times lower in plots where cattle
- 657 were excluded (5.7 ± 1.0 vs. 0.6 ± 0.2 dung piles/ 40 m²). Similarly, in glade edges, cattle+
- buffalo dung density was nearly 7 times lower in plots were cattle were excluded (6.2 ± 1.9
- 659 vs. 0.9 ± 0.4 dung piles/ 40 m²). In plots where megaherbivores were excluded, no
- 660 megaherbivore dung piles were found either inside glades or in glade edges. In plots where
- megaherbivores were allowed, average megaherbivore dung density was 0.9 ± 0.2 dung piles/
- 662 40 m² inside glades and 1.4 ± 0.2 dung piles/ 40 m² in glade edges.

Herbivores alter edge effects Bucher et al.

Highlights

- In east Africa, temporary cattle corrals develop into productive, treeless glades.
- We investigated impacts of different large herbivores on glade edge effects.
- Megaherbivores reduced densities of tall and reproductive trees at glade edges.
- Cattle reduced meso-herbivore use of glade interiors, but not glade edges.
- Large herbivores dampened spatial patterns associated with glades and glade edges.