

Quantifying spatial patterns of grass response to nutrient additions using empirical and neutral semivariogram models

Erica A.H. Smithwick, Douglas C. Baldwin, Kusum J. Naithani

Disturbances influence vegetation patterns at multiple scales, but studies that isolate the effect of scale are rare, meaning that scale and process are often confounded. To explore this, we imposed a large (~3.75 ha) experiment in a South African coastal grassland ecosystem to determine the spatial scale of grass response to nutrient additions. In two of six 60 x 60 m grassland plots, we imposed nutrient additions using a scaled sampling design in which fertilizer was added in replicated sub-plots of varying sizes (1 x 1 m, 2 x 2 m, and 4 x 4 m). The remaining plots either received no additions, or were fertilized evenly across the entire plot area. We calculated empirical semi-variograms for all plots one year following nutrient additions to determine whether the scale of grass response (biomass and nutrient concentrations) corresponded to the scale of the sub-plot additions and compared these results to reference plots (unfertilized or unscaled). In addition, we calculated semi-variograms from a series of simulated landscapes generated using random or structured patterns (neutral models) and compared the semivariogram parameters between simulated and empirical landscapes. Results from the empirical semivariograms showed that there was greater spatial structure in plots that received additions at sub-plot scales, with range values that were closest to the 2 x 2 m grain. These results were in agreement with simulated semivariograms using neutral models, supporting the notion that our empirical results were not confounded by random effects. Overall, our results highlight that neutral models can be combined with empirical semivariograms to identify multi-scalar ecological patterns and this hybrid approach should be used more widely in ecological studies.

1 **Quantifying spatial patterns of grass response to nutrient**
2 **additions using empirical and neutral semivariogram models**

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17 **ABSTRACT**

18 Disturbances influence vegetation patterns at multiple scales, but studies that isolate the effect of
19 scale are rare, meaning that scale and process are often confounded. To explore this, we
20 imposed a large (~3.75 ha) experiment in a South African coastal grassland ecosystem to
21 determine the spatial scale of grass response to nutrient additions. In two of six 60 x 60 m
22 grassland plots, we imposed nutrient additions using a scaled sampling design in which fertilizer
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32 additions at sub-plot scales, with range values that were closest to the 2 x 2 m grain. These
33 results were in agreement with simulated semivariograms using neutral models, supporting the
34 notion that our empirical results were not confounded by random effects. Overall, our results
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36 scalar ecological patterns and this hybrid approach should be used more widely in ecological
37 studies.

38

40 **INTRODUCTION**

41 The importance of spatial autocorrelation for understanding vegetation distributions and
42 processes is well-known (Turner 1989; Watt 1947) and these insights have augmented our
43 understanding of how spatial structure of soils, plants, and climate can regulate ecosystem
44 function, often at multiple, nested scales (Turner et al. 2012). However, for any given study, the
45 scale of this autocorrelation structure and its implications for inferring ecological processes are
46 often not known in advance. Select studies have employed experimental spatial designs *a priori*
47 (Stohlgren et al. 1995) or have used computational models to explore the influence of space on
48 ecosystem properties (Jenerette & Wu 2004; Smithwick et al. 2003; With & Crist 1995).
49 Geostatistical analysis of ecological patterns is also implemented widely (Jackson & Caldwell
50 1993b; Robertson et al. 1993; Smithwick et al. 2005b) but is often used for observations on
51 treatment response rather than to design experimental studies in advance. In the majority of
52 experimental ecological studies, spatial autocorrelation is accounted for *post hoc* through
53 statistical segregation of error terms (Beale et al. 2010).

54 Understanding the scale of ecological processes may help unravel a number of complex
55 ecological questions. For example, disturbances generate structural patterns that may influence
56 ecological processes at many scales (Schoennagel et al. 2008; Turner et al. 2007). However,
57 developing predictive models about the scale of these responses is more difficult because
58 disturbance processes can have cascading effects across scales (Falk et al. 2007; Peters et al.
59 2007). The emergence of bark beetles in the Western U.S. is another multi-scalar process that
60 can only be understood through an unraveling of processes that scale from the tissue-level to the
61 landscape-level (Raffa et al. 2008). Similarly, in savanna science, determining the factors that
62 govern the distribution of grassland and woodland ecosystems is a central focus of many studies
63 (House et al. 2003; Scholes & Archer 1997) and multiple spatial scales are likely needed to

64 explain complex grass-tree interactions (Mills et al. 2006; Okin et al. 2008; Pellegrini 2016;
65 Wang et al. 2010). Herbivore grazing or fire disturbances may each influence savanna nutrient
66 cycling but the scale at which these systems respond is unclear given that the spacing of
67 individual plants, soil properties, terrain, and herbivore assemblages can each contribute to
68 spatial patterns in nutrient availability (Augustine & Frank 2001; Liu et al. 2016; Okin et al.
69 2008; Robertson et al. 1993; Senft et al. 1987; Smithwick et al. 2005b).

70 New approaches are needed to characterize how ecosystems re-organize in response to
71 disturbance and to determine whether there are specific scale or scales of response that are most
72 relevant. In so doing, it may be possible to reduce bias in empirical studies due to scale, thus
73 deepening insights into ecological processes controlling those patterns and minimizing the
74 potential for scale mismatches (Cumming 2011; Ludwig et al. 2000). Implications of such
75 understanding can be significant. For example, Gil et al. (2016) recently found that increasing
76 the scale of nutrient enrichment weakens herbivore control of primary producers, challenging
77 understanding of ecosystem resilience to ecosystem eutrophication. In a more applied setting, it
78 has been shown that the spatial scale of grazer movement can influence nutrient additions in an
79 agricultural field, which could be used to more efficiently target agricultural management
80 activity (Fu et al. 2013). Unraveling such bias or, more optimistically, incorporating scale into
81 ecological prediction and application, requires better experimental approaches to explicitly test
82 for the influence of scale on an ecological pattern.

83 To address this, our overarching goal was to develop a novel methodology to directly test
84 for multi-scalar patterns in primary production following nutrient addition. Our first objective
85 was to determine: at what scale does vegetation respond to nutrient additions? Understanding
86 nutrient limitation to grass productivity is an important question in its own right (Craine et al.

87 2008; Wakeling et al. 2010) but often neglects questions related to the scale of the response. We
88 hypothesized that the grass response would differ between three subplots scales (1 x 1 m, 2 x 2
89 m, and 4 x 4 m). These scales were chosen correspond to ecosystem processes that might govern
90 nutrient uptake, including the spacing of grass tufts, trees or shrubs, and plot-level topography,
91 respectively, which have been identified as critical sources of variation in soil biogeochemistry
92 (Ettema & Wardle 2002; Jackson & Caldwell 1993a; Rietkerk et al. 2000; Smithwick et al.
93 2005b). We surmised that insights gained about the specific scale of response could be used in
94 future studies to understand the scale-specific ecological mechanisms that might be operating at
95 that scale.

96 In addition to testing the within-plot scale of vegetation response to fertilization, our
97 second objective was to test whether there would be stronger spatial structure (enhanced
98 patchiness) across plots in which nutrients were added at multiple sub-plot scales, compared to
99 homogenously fertilized or unfertilized reference plots. To avoid issues of pseudoreplication,
100 these comparisons were made between empirical semivariograms and neutral semivariograms
101 (computer-simulated landscapes that mimic hypothesized patterns). Empirical and neutral
102 semivariograms were compared for six conditions that received, or were artificially modeled to
103 simulate, different nutrient addition patterns (no fertilizer, homogenously fertilized, or fertilized
104 in sub-plot scales) and levels of grazing (fenced or unfenced). We hypothesized that the ranges
105 (autocorrelation distances, or length scales) calculated in the neutral models would be
106 comparable to the ranges calculated from empirical data, and that they would show greater
107 spatial structure in plots that received heterogeneous fertilizers compared to reference plots.

108 In addition, we used these neutral models to explore the influence of herbivory on the
109 spatial structure of fertilized or reference plots. Following nutrient additions, herbivores can

110 influence patterns of primary productivity through enhanced consumption of nutrient-rich
111 patches and/or through enhanced deposition of new nutrients via manure or urine from animals
112 entering the fertilized area, which may have variable effects on spatial heterogeneity depending
113 on the scale of the observations (Fuhlendorf & Smeins 1999). Consumption of nutrient-rich
114 patches may reduce overall variance by reducing differences in biomass amounts compared to
115 ungrazed areas but new nutrient additions by animals also occur when animals move across the
116 plot, or rest in new locations, thus enhancing variability (Auerswald et al. 2010). Therefore, in
117 this study we tested the hypothesis that herbivory increases spatial heterogeneity (Liu et al. 2016)
118 such that patchiness would be highest. i.e., range scales would be smaller, in plots that received
119 both manure additions and scaled nutrient additions (unfenced and heterogeneously fertilized),
120 relative to plots that received only grazing or only homogeneous nutrients.

121 Our third objective was to estimate the degree to which grass biomass was related to
122 patterns in foliar nitrogen content (% N) and phosphorus content (% P), while accounting for
123 random effects due to scale and treatment. Generally, temperate systems are expected to have
124 greater levels of N limitation on vegetation growth than sub-tropical or tropical systems, where P
125 may be more limiting due to highly weathered soils (Domingues et al. 2010; Hedin 2004;
126 Lambers et al. 2008; Vitousek & Sanford 1986). Yet, N, P, and N+P limitations on vegetation
127 productivity have all been documented in African savanna or grassland systems (Augustine et al.
128 2003; Craine et al. 2008; Ngatia et al. 2015; Okin et al. 2008), indicating that an understanding
129 of nutrient limitations on biomass productivity in these systems is still unclear. Complexity in
130 understanding differences in nutrient limitations has been attributed to, for instance, differential
131 nutrient affinities across functional groups (Ratnam et al. 2008; Reich et al. 2003) as well as
132 complex herbivory feedbacks (Augustine et al. 2003), both of which are compounded by issues

133 of scale (Ekblom & Gillson 2010; Fuhlendorf & Smeins 1999). While a complete test of these
134 factors was beyond the scope of this study, we had the opportunity to explore the effect of foliar
135 nutrient concentrations on biomass responses under a dual nutrient addition experiment, while
136 treating the levels of our experimental design (e.g. plot, subplot, treatment) as separate random
137 effects. By so doing, we were able to explore whether foliar N, P or N and P together explained
138 patterns in biomass, beyond the given variability in our study design, and to determine the level
139 (scale) at which these random effects were most pronounced.

140

141 **METHODS**

142 **Study area.** This study was conducted in Mkambathi Nature Reserve, a 7720-ha protected area
143 located at 31° 13' 27" S and 29° 57' 58" E along the Wild Coast region of the Eastern Cape
144 Province, South Africa. The Eastern Cape is at the confluence of four major vegetative
145 groupings (Afromontane, Cape, Tongaland-Pondoland, and Karoo-Namib) reflecting
146 biogeographically complex evolutionary histories. It is located within the Maputaland-
147 Pondoland-Albany conservation area, which bridges the coastal forests of Eastern Africa to the
148 north, and the Cape Floristic Region and Succulent Karoo to the south and west. The
149 Maputaland-Pondoland-Albany region is the second richest floristic region in Africa, with over
150 8,100 species identified (23 % endemic), and 1,524 vascular plant genera (39 endemic).
151 Vegetation in Mkambathi is dominated by coastal sour grassveld ecosystems, which dominate
152 about 80 % of the ecosystem (Kerley et al. 1995; Shackleton et al. 1991), with small pockets of
153 forest along river gorges, wetland depressions, and coastal dunes. Dominant grasses in the
154 Mkambathi reserve include the coastal *Themeda triandra* – *Centella asiatica* grass community,
155 the tall grass *Cymbopogon validus* – *Digitaria natalensis* community in drier locations, and the

156 short-grass *Tristachya leucothrix-Loudetia simplex* community (Shackleton 1990). Grasslands in
157 Mkambathi have high fire frequencies, and typically burn biennially. Soils are generally derived
158 from weathered Natal Group sandstone and are highly acidic and sandy with weak structure and
159 soil moisture holding capacity (Shackleton et al. 1991). Climate is generally considered warm
160 temperate, where average monthly minimum and maximum temperature in East London is 14
161 and 23 °C respectively, and precipitation is 921 mm yr⁻¹ (1961 – 1990 average, S.A. Weather
162 Service).

163 We established a large-scale experimental site that included six 60 x 60 m plots arranged
164 in a rectangular grid (Eastern Cape Parks and Tourism Agency Permit RA0081). The site was
165 surrounded by a fuel-removal fire-break and each plot was separated by at least 10 m for a total
166 size of 3.75 ha for the entire site. To account for grazing, a fence was constructed around three
167 of these plots.

168 **Nutrient additions.** Nutrient additions were applied to four plots whereas two plots received no
169 fertilizer additions. Of the four plots that received fertilization, two received nutrients evenly
170 across the entire 60 x 60 m plot (“homogenous plots”) and the other two fertilized plots received
171 nutrient additions within smaller subplots in a heterogeneous design (“heterogeneous plots”).
172 Within heterogeneous plots, fertilizer was applied within subplots of three different sizes (1 x 1
173 m, 2 x 2 m, and 4 x 4 m) that were replicated randomly across each plot (**Fig. 1**). Location of
174 individual subplots was determined prior to field work using a Latin Hypercube random
175 generator that optimizes the variability of lag distances among sampling plots and is ideal for
176 geostatistical analysis (Xu et al. 2005). There were a total of 126 subplots per plot that received
177 fertilizer in the heterogeneous plots. All sampling locations were geo-referenced with a GPS

178 (Trimble). The number of sub-plot units at each scale was determined so as to equalize the total
179 fertilized area at each sub-plot scale (i.e., six 4 x 4 m plots and 24, 2 x 2 m plots).

180 To ensure aboveground grass biomass would respond to nutrient additions, we employed
181 a dual (nitrogen (N) + phosphorus (P)) nutrient addition experiment. Additional N was added as
182 either ammonium nitrate (230 g kg⁻¹ N) or urea (460 g kg⁻¹) at a rate of 10 g m⁻² yr⁻¹ in a single
183 application, following the protocols of Craine et al. (2008). Additional P was added as
184 superphosphate (105 g kg⁻¹ P) at a rate of 5 g m⁻² yr⁻¹. Dual addition (N+P) was chosen to
185 increase the likelihood of treatment response and increase geostatistical power by reducing the
186 number of treatments, thus increasing sample size. Per unit area, levels of nutrient additions
187 were constant among plots and subplots.

188 **Vegetation and Soil Sampling.** One year following nutrient additions, a subset of subplots was
189 sampled for soil and vegetation nutrient concentrations and biomass. Sub-plots to be sampled
190 were randomly selected prior to being in the field using the Latin Hypercube approach. The
191 approach allowed us to specify a balanced selection of subplots within each subplot size class
192 (four 4 x 4 m, eight 2 x 2 m, and thirty-two 1 x 1 m). Within each subplot location that was
193 revisited, we randomly selected locations for biomass measurement and vegetation clippings.
194 Two locations were identified and flagged from within the 1 x 1 m subplots (center coordinate
195 and a random location 0.5 m from center), four samples were identified and flagged from within
196 the 2 x 2 m subplots, and eight samples were identified and flagged from within the 4 x 4 m
197 subplots. At each flagged location within sampled subplots, productivity was measured as grass
198 biomass using a disc pasture meter (DPM; Bransby and Tainton, 1977) and grab samples of grass
199 clippings were collected for foliar nutrient analysis, using shears and cutting to ground-level.
200 Calibration of the DPM readings was determined using ten random 1 x 1 m subplots in each plot

201 (n = 60 total) that were not used for vegetation or soil harvesting, in which the entire biomass
202 was harvested to bare soil. Soil samples from the top 0 – 10 cm soil profile depth were collected
203 adjacent to vegetation samples. Due to logistical and financial constraints, these samples were
204 collected in fenced plots only. The A horizon of the Mollisols was consistently thicker than 10
205 cm, so all samples collected were drawn from the A horizon. Soil samples were shipped to
206 BEMLab (Strand, South Africa) for nutrient analysis.

207 **Laboratory Analysis.** Biomass samples were separated into grasses and forbs, weighed, dried
208 for 24 h at 60 °C, and reweighed. Vegetation nutrient samples were dried, ground with a 40 mm
209 grinding mesh, and then shipped to the Penn State Agricultural Analytical Laboratory
210 (University Park, Pennsylvania; USDA Permit PDEP11-00029). Grass P concentration was
211 analyzed using a hot block acid digestion approach (Huang & Schulte 1985) and grass N
212 concentration was measured with a Combustion-Elementar Vario Max method (Horneck &
213 Miller 1998). Soil N and C concentrations were determined on a LECO elemental analyzer
214 (Leco Corporation, St. Joseph, MI). Soil P was analyzed using acid extraction following the
215 method of Wolf and Beegle (1995). Soil pH was estimated using KCl extraction following
216 Eckert and Sims (1995).

217

218 **Calculations and Statistics.** Aboveground grass biomass was estimated from DPM
219 measurements, using linear regression to relate DPM estimates with harvested biomass at
220 calibration subplots. DPM estimates were significantly correlated with total grass biomass ($R^2 =$
221 0.76, $p < 0.0001$) (**Supplementary Material, Fig S1**) and the resulting equation was then used to
222 estimate biomass at the remaining 606 locations.

223 To test the scale of grass biomass response to fertilization (Objective 1), we used a
 224 maximum likelihood approach to quantify the spatial structure of biomass separately for each of
 225 the six empirical plots. The approach assumes that the data ($Y_1 \dots Y_n$) are realizations of an
 226 underlying spatial process, and that the distribution of the data follows a Gaussian multivariate
 227 distribution:

$$228 \quad Y \sim N(\mu \mathbf{1}, C \Sigma + C_0 I) \quad (1)$$

229 where μ is the mean of the data multiplied by an n-dimensional vector of 1's, C is the partial sill
 230 (total sill = $C_0 + C$), Σ is an n x n spatial covariance matrix, C_0 is the nugget effect, and I is an n
 231 x n identity matrix. The i,j th element of Σ is calculated with a spatial covariance function $\rho(h_{ij})$,
 232 where h_{ij} is the Euclidean distance between measurement points i and j . An exponential
 233 covariance model was chosen for its relative simplicity. The full equation for summarizing the
 234 second order moment for an element i,j is:

$$235 \quad \gamma(h_{ij}) = C_0 + C \left[\exp\left(\frac{-h_{ij}}{\phi}\right) \right] \quad (2)$$

236 where $\gamma(h_{ij})$ is the modeled spatial covariance for measurements i and j , ϕ is the range parameter,
 237 and $3 * \phi$ is the range of spatial autocorrelation. The underlying spatial mean μ may be held
 238 constant or estimated with a linear model across all locations. We used the plot-level mean of
 239 the data for μ (Table 1).

240 The measured soil and plant variables exhibited varying degrees of non-normality in their
 241 distributions, which violates the assumption of Gaussian stationarity within the underlying
 242 spatial data generating process. To uphold this assumption, we transformed variables at each
 243 plot using a box-cox transformation (Box & Cox 1964):

$$244 \quad \hat{Y}_i = (Y_i^\lambda - 1) / \lambda \quad \text{if } \lambda \neq 0 \quad (3)$$

$$245 \quad \hat{Y}_i = \log(Y_i) \quad \text{if } \lambda = 0$$

246 where Y_i is an untransformed variable (e.g., biomass) at location i , \hat{Y}_i is the transformed variable,
247 and λ is a transformation parameter. We optimized the three spatial covariance model
248 parameters and the transformation parameter (C_0, C, ϕ, λ) with the maximum likelihood
249 procedure. A numerical finite-difference approximation algorithm selected the set of parameters
250 that maximized a normal multivariate log-likelihood function (more details in Diggle et al,
251 2003). To approximate a sampling distribution of each parameter, a bootstrapping algorithm was
252 used where a randomly sampled subset of data was inputted into the same maximum likelihood
253 approach for 1000 iterations. This provided a population of fitted parameters and models that
254 was used to analyze the approximate distributions of each parameter for each plot. The
255 maximum likelihood optimization was cross-validated by removing a random sub-sample of
256 measurements from the optimization and then using the optimized model to make predictions at
257 locations where measurements were removed. Observed vs. predicted values from the cross-
258 validation procedure were then analyzed at each plot separately.

259 We used ordinary kriging (Cressie 1988) with the optimized spatial covariance model
260 from the maximum likelihood analysis to estimate biomass across all plots. Ordinary kriging is
261 useful in this case, because we detected spatial structure in the biomass data when considering all
262 biomass data at once (see Results). The geoR package (Ribeiro Jr. & Diggle 2001) in the R
263 statistical language (R Development Team, 2014) was used for all spatial modeling and kriging.

264 To compare spatial structure across empirical or simulated patterns (Objective 2), we
265 compared the empirical semivariograms with neutral semivariogram models. Essentially this
266 approach allows us to compare empirical patterns across a set of null models in which the
267 patterns are known. The neutral semivariogram models were constructed for six simulated

268 landscapes (**Fig. 2**) to represent alternative landscape structures in response to nutrient addition
269 and grazing: **(a)** fenced-unfertilized (biomass was assumed to be randomly distributed around the
270 mean of the biomass from the fenced, unfertilized experimental plot), **(b)** fenced-heterogeneous
271 (biomass of **(a)** was doubled for selected subplots, following the same subplot structure that was
272 used in the field experiments), **(c)** fenced-homogenous (biomass of **(a)** was doubled at every grid
273 cell to mimic an evenly distributed fertilization response), **(d)** unfenced-unfertilized (biomass of
274 **(a)** was increased by 50 % in response to a combined effect of biomass loss by grazing and
275 biomass gain by manure nutrient additions by herbivores; the increase occurred at a subset of
276 sites to mimic random movement patterns of herbivores), **(e)** unfenced-heterogeneous (biomass
277 equaled biomass of herbivory only, fertilizer only, or herbivory + fertilizer), and **(f)** unfenced-
278 homogenous (biomass of **(d)** was doubled at all grid cells to mimic the additive effects of
279 herbivores and homogenous fertilizer additions).

280 The spatial structure of simulated landscapes was analyzed using the same maximum
281 likelihood approach as described above and data was not transformed. The mean (μ) was
282 estimated using a constant trend estimate. To compare neutral (simulated data) and empirical
283 (observed data) semivariogram models, we scaled the nugget and sill model parameters by
284 dividing these parameters into the maximum observed semivariance value within each plot.

285 The relationship between biomass and foliar N and P (Objective 3) was investigated
286 using a linear mixed modeling approach, where herbivory, fertilizer type (i.e., heterogeneous,
287 homogenous, and unfertilized), plot treatment, and subplot size were all included as random
288 effects. Accounting for experimental factors as random effects is a desirable approach for
289 managing non-independence in data collected across different treatments that are not replicated
290 (i.e., pseudoreplication) (Millar & Anderson 2004). Multiple combinations of random effects

291 and fixed effects were tested, where foliar N and P represented fixed effects upon biomass, and
292 model error was assumed to be Gaussian. A normal likelihood function was minimized to
293 estimate optimal regression coefficients for each mixed model formulation.

294 Our objective was to identify a mixed model that estimated biomass closely to
295 observations, while also having the fewest possible parameters. The Akaike's Information
296 Criterion (AIC) and Bayesian Information criterion (BIC) were used to compare different
297 models, since they decrease with a negative log-likelihood function but increase with the amount
298 of parameters used in the model (Burnham and Anderson, 2004). After all model formulations
299 were fitted to observed biomass and both AIC and BIC were calculated for each model, we chose
300 the model with the lowest BIC as best representing the tradeoff of parsimony and prediction
301 skill. The BIC associated with all other models was subtracted into the lowest available BIC,
302 and models with a difference in $BIC > 2$ were deemed significantly less favorable at estimating
303 biomass and representing random effects than the model with the lowest BIC. All mixed
304 modeling was conducted with the R package lme4.

305

306 RESULTS

307 Across plots, vegetation biomass ranged from $376 \pm 6.0 \text{ g m}^{-2}$ (mean \pm 1 Standard Error
308 (SE), $n = 124$) in the unfenced, homogeneously fertilized plot to $563 \pm 18.6 \text{ g m}^{-2}$ ($n = 128$) in the
309 unfenced, heterogeneously fertilized plot (**Table 1**). Forb biomass was approximately 18 % of
310 total biomass, with the remaining 82 % represented by grass. Vegetation nutrient concentrations
311 increased, and N:P ratios declined, following fertilization (**Table 1**). Vegetation N concentration
312 ranged from $0.58 \pm 0.01 \%$ to $0.75 \pm 0.04 \%$, averaging $0.60 \pm 0.01 \%$ in unfertilized plots versus
313 an average of $0.72 \pm 0.02 \%$ in heterogeneously fertilized plots and $0.77 \pm 0.02 \%$ in

314 homogeneously fertilized plots, an increase of 20 % and 28 %, respectively. Vegetation P
315 concentration averaged $0.037 \pm 0.001 \text{ mg g}^{-1}$ in unfertilized plots, $0.056 \pm 0.002 \text{ mg g}^{-1}$ in
316 heterogeneously fertilized, and $0.057 \pm 0.002 \text{ mg g}^{-1}$ in homogeneously fertilized plots, an
317 increase of 34 and 35%, respectively. The N:P ratios ranged from a high of 17.9 in the fenced-
318 unfertilized plot to 12.1 in the unfenced-homogeneously fertilized plot. Vegetation C content
319 averaged $44.6 \pm 0.13 \%$ across all six plots. Soil P and N were also higher following fertilization
320 in the fenced plots, where these variables were measured (**Supplementary Table 1**). Soil C
321 ranged from $2.49 \pm 0.01 \%$ to $2.55 \pm 0.01 \%$ across plots. Soil pH was 4.27 in the unfertilized
322 plot and 4.08 in fertilized plots. Confirming reference conditions, pH measured in a single
323 control plot in 2011 prior to fertilization was 4.21 ± 0.01 .

324 We had expected that biomass and vegetation nutrient concentrations following
325 fertilization would be spatially autocorrelated at scales corresponding to the scale of the
326 fertilization additions (Objective 1). In heterogeneously fertilized plots this would be indicated
327 by range distances from empirical semivariograms that corresponded to the hypotenuse distances
328 of the subplot scales (i.e., 1 m, 2.83 m, and 5.66 m hypotenuse distances for the 1 x 1 m, 2 x 2 m,
329 and 4 x 4 m subplots, respectively). In homogeneously fertilized plots or unfertilized plots, spatial
330 structure would be observed at scales other than scales of the subplots (or not at all) and we
331 would expect a higher degree of nugget variance (spatial structure expressed at scales finer than
332 those that were measured). Results from the empirical semi-variogram model of the
333 heterogeneous plots show that there was a statistically significant patch structure at scales similar
334 to the scale of the subplots in both of the heterogeneously fertilized plots (**Fig. 3b,f**). Also
335 confirming these expectations, in unfenced and fenced plots (unfertilized or homogeneously
336 fertilized) the range scale was significantly longer or shorter (**Fig. 3; Supplementary Table 2**).

337 Examining the sampling distributions of the semivariogram range values for vegetation
338 biomass determined from the maximum likelihood and bootstrapping analysis, it can be seen that
339 the range value most closely resembles that of the hypotenuse of the 2 x 2 m subplot, relative to
340 the other subplots (**Fig. 3d,h**). Higher spatial structure in the heterogeneous versus
341 homogeneous or unfertilized plots can also be seen in the kriged plots of biomass (**Fig. 4**). These
342 maps also demonstrated the higher mean levels of biomass in fertilized subplots relative to areas
343 outside of subplots or relative to other plots. Normalized nugget/sill ratios were highest in the
344 unfenced, homogeneously fertilized plot (3.89) with lower ratios (0-0.02) for heterogeneously
345 fertilized or fenced treatments.

346 Consistent with the biomass results, semivariogram parameters for vegetation % N and %
347 P (**Supplementary Table 3**) generally supported expectations, at least for the heterogeneously
348 fertilized and fenced plot, in which range scales were comparable to subplot scales in the (% P, ~
349 4.9 m, % N, ~ 5.8 m); however, other plots showed higher or lower ranges. The proportion of
350 nugget variance in semivariogram models of vegetation % N and % P was highest in the
351 unfertilized plots, which supports expectations of higher spatial structure in fertilized treatments,
352 as for biomass, but differences were variable among individual plots. However, semi-variograms
353 of soil carbon and nutrients (**Supplementary Table 3**) showed few differences among model
354 parameters among treatments where these were measured (fenced plots, only). In sum,
355 differences in spatial structure that could be attributed to treatments were most evident for
356 biomass, somewhat evident for vegetation % N and % P, and least evident for soil C or nutrients.

357 To compare mean differences among plots (Objective 2), we compared normalized
358 spatial model fits of neutral and empirical semi-variograms, which allowed us to quantify the
359 relative influence of treatment responses to random processes not accounted for in the

360 experimental design. Results indicated that the range of autocorrelation in heterogeneous plots
361 was similar to that found in the empirical semi-variograms and generally matched subplot scales
362 (**Fig. 5**). Interestingly, range scales were higher, and proportional nugget variance was greater in
363 fenced plots, compared to unfenced plots. This was an opposite pattern than that observed in
364 empirical plots, in which grazing resulted in longer range scales and higher nugget variance.
365 These results likely reflect processes not included in the neutral model. Particularly, the neutral
366 model was designed to mimic patterns in biomass using assumptions about herbivore movement,
367 grazing, and manure additions, assuming they were additive to the effects of fertilization,
368 whereas empirical results likely reflect complex interactions between grazing and fertilization.
369 In this case, longer observed range scales in grazed versus ungrazed plots may reflect enhanced
370 homogenization of biomass through grazing, an effect that was apparently greater than biomass
371 enhancement from nutrient addition. Nonetheless, the fact that heterogeneous plots were similar
372 between fenced and unfenced plots indicates the strong scalar influence of nutrient additions
373 relative to nutrient-herbivore interactions.

374 Addressing Objective 3, mixed models used to estimate biomass from N or P foliar
375 concentrations, while treating plot and treatment as random effects, showed that biomass was
376 best predicted by levels of foliar P, relative to foliar N alone or foliar N x P (**Table 2**). Although
377 foliar P alone did better than foliar N alone as a fixed effect, the difference was marginal (< 2
378 BIC). The ‘best’ model uses only plot treatment type as a random effect, which outperformed
379 model formulations using herbivory or fertilizer type and those with nested structures
380 incorporating subplot size as random effects.

381

382 **DISCUSSION**

383 Scale issues confound a complete understanding of ecosystem response to perturbations
384 (Turner 2010). This is particularly important in grassland or savanna systems in which the
385 relative importance of factors affecting grass-woodland ecosystems are debated (e.g., Mills et al.
386 2006; Sankaran et al. 2004; Scholes & Archer 1997), with some suggesting that these factors are
387 nested hierarchically with spatial scale (Pellegrini 2016; Pickett et al. 2003; Rogers 2003).
388 Results of this experiment yield data on the spatial scale of the nutrient-productivity relationship
389 in a grassland coastal forest of the Eastern Cape, South Africa, and support the assertion that
390 ecological processes are multi-scaled and hierarchical in response to nutrient additions.
391 Specifically, we observed immediate (one year following fertilization) biomass response at the
392 plot level due to nutrient addition and fencing. Using spatial modeling, we discerned strong
393 autocorrelation structure in biomass at subplot scales associated with scales of nutrient addition.
394 The strongest evidence for spatial structure was observed at the 2 x 2 m scale. Comparing this
395 spatial structure between neutral and empirical semivariogram models, we were able to infer the
396 relative importance of treatment conditions (nutrients and grazing) to random, spatially
397 structured processes. Traditional ecosystem-level approaches to quantify spatial pattern have
398 focused on understanding extant patterns, in which processes are necessarily inferred or
399 correlated. However, by examining spatial patterns under experimental conditions in which scale
400 and treatment are controlled, we were able to relate ecological responses to known scales of
401 perturbation.

402 Subplots responded very strongly to nutrient additions at subplot scales, as evident from
403 the kriged maps, showing that most of the increase in average plot biomass was due to large
404 increases in subplot biomass. One surprising result of our study was the detection of average
405 greater biomass in plots that received heterogeneous fertilizer applications compared to plots that

406 received homogeneous fertilizer, despite the fact that fertilizer was added equally on a per area
407 basis. Several other studies have found higher biomass following heterogeneous nutrient
408 applications. For example, Day et al. (2003) observed that heterogeneous spatial patterns of
409 nutrient supply in early stages of grassland development led to enhanced nutrient acquisition and
410 biomass productivity. Similarly, Du et al. (2012) observed increased plant biomass following
411 heterogeneous nutrient fertilization in old-field communities in China. Mechanisms for
412 enhanced productivity following heterogeneous nutrient supply are not clear but may include
413 shifts in root structure and function or shifts in species dominance, which were not analyzed
414 here. For example, roots may respond to patchiness in nutrient availability by modifying root
415 lifespan, rooting structures and uptake rate to maximize nutrient supply (Hodge 2004; Robinson
416 1994). In turn, initial advantages afforded by plants in nutrient-rich locations may result in
417 larger plants and advantages against competitive species, potentially via enhanced root growth
418 (Casper et al. 2000).

419 Observing vegetative response to nutrient additions is complicated by processes such as
420 luxury consumption (Ostertag 2010), initial spatial patterns in soil fertility (Castrignano et al.
421 2000), root distribution, signaling and allocation (Aiken & Smucker 1996), species and
422 functional group shifts (Ratnam et al. 2008; Reich et al. 2003), or species' differences in uptake
423 rates or resorption (Reed et al. 2012; Townsend et al. 2007). Spatial patterns of finer-scale
424 processes such as microbial community composition have also been explored and are known to
425 influence rates of nutrient cycling (Ritz et al. 2004; Smithwick et al. 2005a). In the case of
426 heterogeneous nutrient supply, species competitive relationships across space may be enhanced
427 (Du et al. 2012) and may result in increases in plant diversity (Fitter 1982; Wijesinghe et al.
428 2005), although other studies have found little evidence to support this claim (Gundale et al.

429 2011). Together, these factors may explain the high unexplained variance of vegetation N and P
430 concentrations that we observed and may account for the nugget variance in our empirical spatial
431 models. However, effects of species composition shifts on plant biomass were likely minimal in
432 this study given the short-term nature of the study (one year), although the patchiness in biomass
433 in gridded fertilizer plots indicates size differences that are likely to modify competitive
434 relationships in the future. Unfortunately, the site burned one year following the experiment,
435 precluding additional tests of these relationships.

436 Understanding the length scales of key ecosystem properties is critical for determining
437 optimal scales for studying ecological systems, interpreting change in ecological communities,
438 and assessing landscape connectivity and ecosystem resilience (Johnson 2009; Turner et al.
439 2012). Autocorrelation structure may reflect influences of spatial interactions among
440 individuals, patches, or abiotic gradients (Ettema & Wardle 2002; Jackson & Caldwell 1993a;
441 Rietkerk et al. 2000; Smithwick et al. 2005b). Our study showed that biomass responded
442 strongly at all scales in which nutrients were added, but was strongest at the 2 x 2 m scale.
443 Rietkerk et al. (2000) observed patchiness in soil moisture at three unique scales (0.5 m, 1.8 m
444 and 2.8 m) in response to herbivore impacts. Following fire in the Greater Yellowstone
445 Ecosystem (Wyoming, U.S.A.), Turner et al. (2011) observed variation in soil properties at the
446 level of individual soil cores, and Smithwick et al. (2012) observed autocorrelation in post-fire
447 soil microbial variables that ranged from 1.5 to 10.5 m. In savanna and boreal ecosystems,
448 patchiness in soil resources at the level of individual shrubs and trees has been demonstrated by
449 several studies (Dijkstra et al. 2006; Hibbard et al. 2001; Lechmere-Oertel et al. 2005; Liski
450 1995; Pennanen et al. 1999). Obviously, other contingent factors (e.g., herbivory, historical fire,
451 land use) are important for explaining grass species distributions and productivity patterns at

452 broader temporal and spatial scales, but our study suggests that intermediate length scales (<4
453 m²) reflect ecological processes important for understanding patterns in grassland nutrition.

454 Many studies have used stoichiometric relationships of N and P to infer nutrient
455 limitation (Koerselman & Meuleman 1996; Reich & Oleksyn 2004), although there are limits to
456 this approach (Ostertag 2010; Townsend et al. 2007). Using this index, our N:P ratios of
457 vegetation in reference plots would indicate co-limitation for N and P prior to fertilization (N:P >
458 16). Addition of dual fertilizer appeared to alleviate P limitation more than N, with N:P ratios
459 reduced one year following treatment, indicating N limitation or co-limitation with another
460 element (N:P < 14). In addition, P was most strongly correlated with biomass in linear mixed
461 effects modeling across all plots, after accounting for random effects of the experimental design.
462 Ostertag (2010) also showed that there was a preference for P uptake in a nutrient limited
463 ecosystem in Hawaii and suggested that foliar P accumulation may be a strategy to cope with
464 variability in P availability. Grazing may also preferentially increase grass P concentrations in
465 semi-arid systems in South Africa (Mbatha & Ward 2010). The cumulative impacts of
466 preferential plant P uptake and P additions from manure may explain the high spatial structure
467 observed in our grazed and fertilized plots.

468 Spatial heterogeneity may be critical for influencing ecosystem resilience to stressors by
469 mediating disturbance spread (i.e., fire, invasive species dispersal) (Yang et al. 2008), explaining
470 patterns in disease emergence (Wu et al. 2015), or promoting recovery and regeneration through
471 maintenance of legacy structures and functions (Turner et al. 2012). Understanding the
472 importance of spatial pattern for maintaining soil fertility is increasingly relevant for agricultural
473 management that incorporates precision technology as well as for ecosystem management
474 activities that prioritize monitoring and restoration. In South Africa, grasslands cover nearly

475 one-third of the country and maintain the second-highest levels of biodiversity, behind the Cape
476 Floristic Region. Grasslands of Mkambati are representative of ancient grasslands that are under
477 threat globally, yet possess high levels of species endemism and provide significant ecosystem
478 goods and services (Bond 2016). Grasslands in southern Africa are expected to undergo
479 significant losses in biodiversity in coming decades due to increasing pressure from agricultural
480 development and direct changes in climate (Biggs et al. 2008; Huntley & Barnard 2012). For
481 example, losses in critical African bird species in fynbos and grassland biomass are expected to
482 be 30 – 40 % lower by 2085 as a result of projected climate change, with range reductions > 60
483 % in many cases. As a result of these pressures, understanding the factors that regulate
484 ecosystem productivity, and the scales at which they operate, is critical for guiding ecosystem
485 management activities aimed at maintaining landscape sustainability.

486

487 **CONCLUSIONS**

488 This study provided an opportunity for direct hypothesis-testing of the influence of
489 spatial pattern on ecosystem processes, providing an alternative to studies in which spatial
490 autocorrelation is observed post-hoc or otherwise confounds complex ecosystem dynamics.
491 Specifically, spatial models were used to test the scale at which vegetation responded to nutrient
492 additions within plots and we compared these model parameters to those determined from model
493 parameters from simulated landscapes using replicated neutral models. Biomass responded to
494 nutrient additions, with spatial autocorrelation of the biomass response highest at the 2 x 2 m
495 scale. Comparison of empirical and simulated neutral models, confirmed that this length scale
496 accounted for the majority of the spatial structure in the observations. Vegetation P
497 concentrations were elevated more than vegetation N suggesting plant P preference or luxury

498 consumption, although vegetation nutrient concentrations exhibited lower spatial structure than
499 that observed for biomass. Finally, this study was undertaken in a nature reserve in which
500 grasslands are being actively managed to meet multiple management objectives including
501 wildlife management and biodiversity. These results indicate strong coupling between soil
502 nutrients and aboveground function which should be considered in vulnerability assessments that
503 consider grassland responses to shifts in climate and management activities.

504

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Table 1 (on next page)

Plot-level biomass and vegetation nutrient concentrations.

Mean (± 1 standard error (SE)) biomass, vegetation N concentration, vegetation P concentration, and N:P ratios across experimental plots in Mkambathi Nature Reserve, one year following nutrient fertilization.

1

Treatment	Average Biomass ± 1 SE (g m⁻²)	Average N ± 1 SE (%)	Average P ± 1 SE (%)	N:P	n
Fenced Unfertilized	411.9 ± 9.75	0.646 ± 0.024	0.036 ± 0.001	17.9	134
Heterogeneous	542.4 ± 15.05	0.747 ± 0.041	0.048 ± 0.002	15.6	120
Homogeneous	456.2 ± 8.28	0.710 ± 0.014	0.054 ± 0.002	13.2	117
Unfenced Unfertilized	483.6 ± 13.70	0.576 ± 0.011	0.038 ± 0.001	15.2	132
Heterogeneous	562.6 ± 18.60	0.775 ± 0.015	0.064 ± 0.002	12.1	128
Homogeneous	375.4 ± 5.96	0.722 ± 0.017	0.059 ± 0.002	12.2	124

2

Table 2 (on next page)

Mixed model results comparing biomass to foliar nutrients.

Results of the mixed model relating biomass to foliar nutrients, where herbivory, fertilizer type, plot treatment, and subplot size were all tested as random effects; foliar N and P represented fixed effects upon biomass, and model error was assumed to be Gaussian. A normal likelihood function was minimized to estimate optimal regression coefficients for each mixed model formulation. Both Akaike's Information Criterion (AIC) and Bayesian Information criterion (BIC) were used to compare different models. Delta (Δ) represents differences in BIC between the current model and the model with the lowest BIC.

1
2

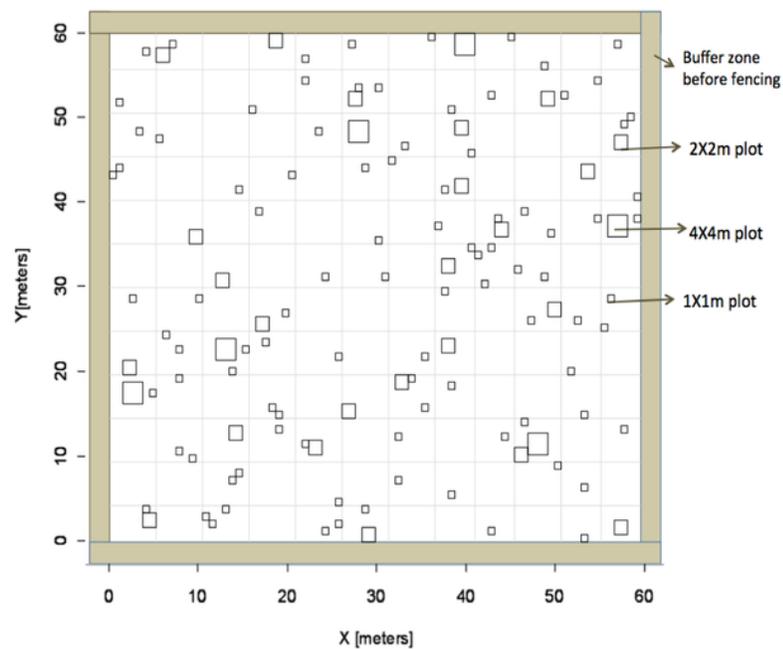
3

Model	DF	AIC	BIC	Δ
<i>Random Effects</i>				
Plot	5	1092.4	1114.2	0.0
Herbivore	5	1190.1	1211.9	97.7
Fertilizer	5	1100.7	1122.5	8.3
Plot Sub-Plot	6	1090.4	1116.5	2.3
Herbivore Sub-Plot	6	1188.6	1214.7	100.5
Fertilizer Sub-Plot	6	1102.7	1128.8	14.6
<i>Fixed Effects</i>				
N + P	5	1090.3	1112.1	5.3
P	4	1089.8	1107.3	0.4
N	4	1090.7	1108.2	1.3
N : P	6	1092.3	1118.5	11.6
N + P + Sub-Plot	6	1092.3	1118.5	11.6
N + P : Sub-Plot	8	1095.6	1130.5	23.6
P + N ²	5	1091.6	1113.4	6.6
N + P ²	5	1089.7	1111.5	4.7
N ² + P ²	5	1091.1	1113.0	6.1
N ²	4	1093.3	1110.8	3.9
P²	4	1089.4	1106.9	0.0

1

Experimental design

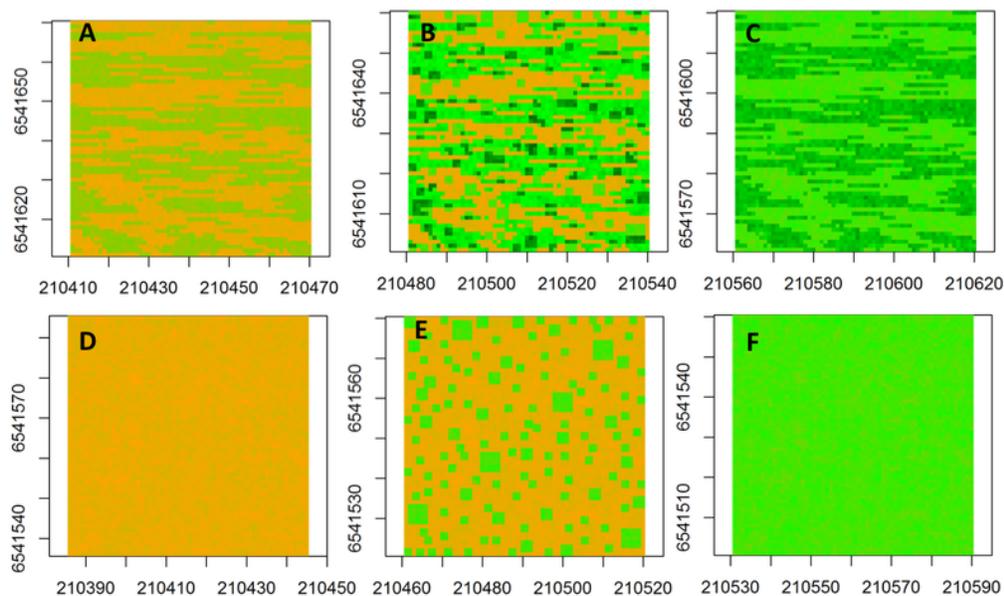
Overview of experimental design based on Latin Hypercube sampling used to identify subplot locations to receive fertilizer in the heterogeneous plots.



2

Spatial maps of neutral models

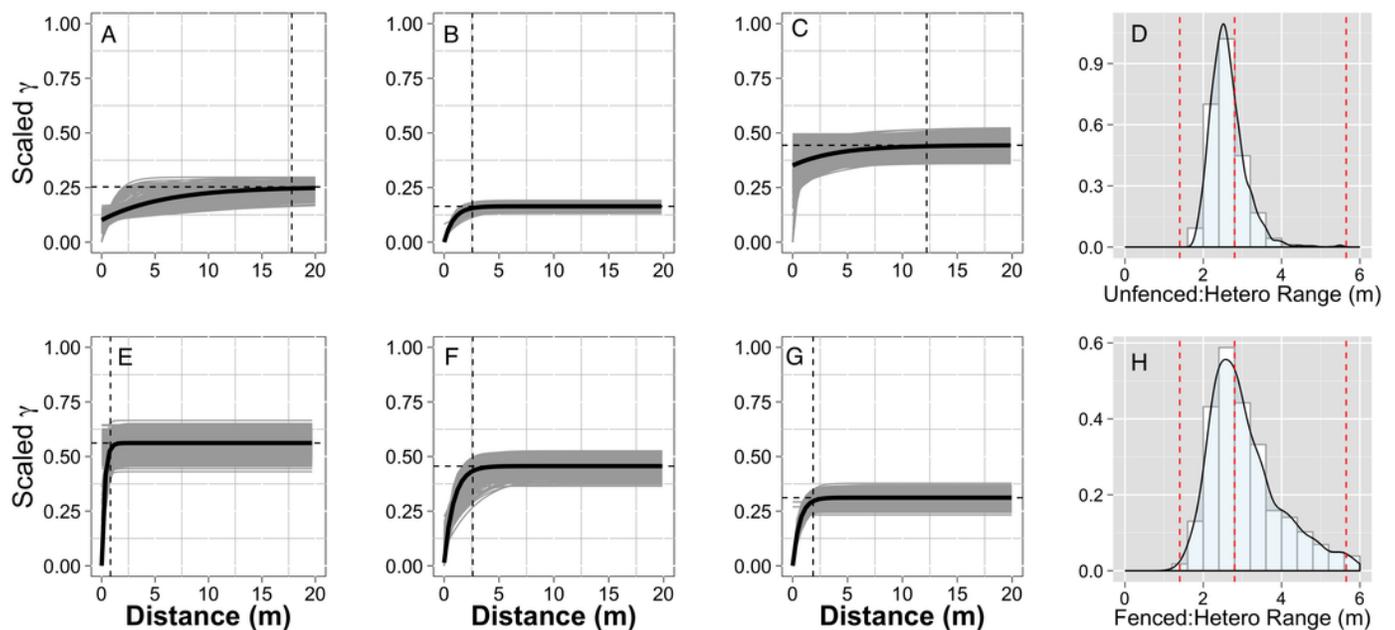
Spatial maps of neutral models used to simulate vegetation biomass for the following conditions: (a) unfenced, unfertilized, (b) Unfenced, heterogeneously fertilized, (c) Unfenced-homogeneously fertilized, (d) Fenced, unfertilized, (e) Fenced, heterogeneously fertilized, (f) Fenced, homogeneously fertilized.



3

Empirical semivariograms

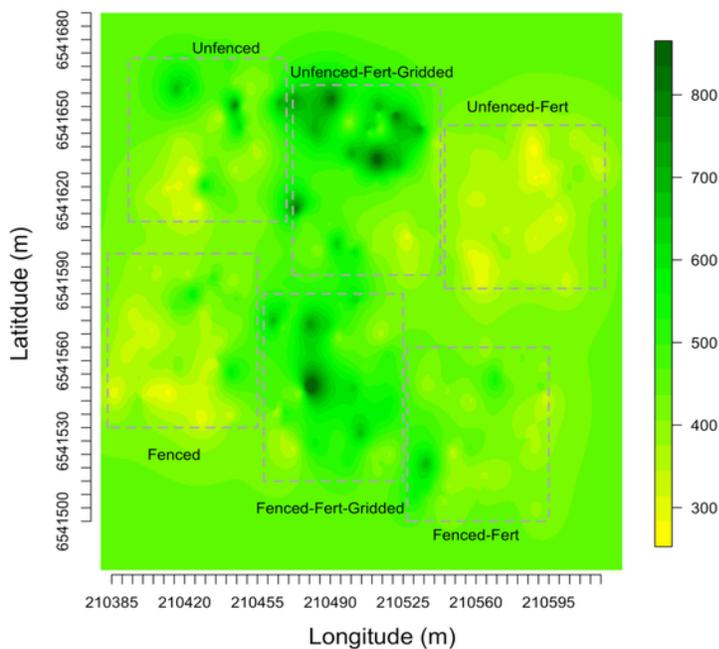
Empirical semi-variograms of vegetation biomass for each plot: (A) Unfenced, unfertilized, (B) Unfenced, Heterogeneously Fertilized, (C) Unfenced, homogeneously fertilized, (E) Fenced, unfertilized, (f) Fenced, heterogeneously fertilized, (G) Fenced, homogeneously fertilized. Shaded lines represent semi-variogram models fitted during the bootstrapping procedure. Dashed vertical line represents the range value. Also shown: the sampling distribution of the range parameter for heterogeneously fertilized plots that were either (D) Unfenced, or (H) Fenced. The distribution was calculated with a bootstrapping approach with maximum likelihood optimization. Dashed vertical lines represent the hypotenuses of the 1x1 m (1.4), 2x2 (2.8), and 4x4 (5.7) sub-plots.



4

Kriged biomass map.

Kriged map of biomass using ordinary kriging with a spatial covariance model optimized by a maximum likelihood analysis: (A) Unfenced, unfertilized, (B) Unfenced, heterogeneously fertilized, (C) Unfenced, homogeneously fertilized, (D) Fenced, unfertilized, (E) Fenced-heterogeneously fertilized, (F) Fenced, homogeneously fertilized.



5

Semivariograms from neutral models.

Simulated semivariograms of vegetation biomass for each plot from neutral landscape models: (A) Unfenced, unfertilized, (B) Unfenced, heterogeneously fertilized, (C) Unfenced, homogeneously fertilized, (D) Fenced, unfertilized, (E) Fenced, heterogeneously fertilized, (F) Fenced, homogeneously fertilized. Shaded lines represent semi-variogram models fitted during the bootstrapping procedure. Dashed vertical line represents the optimal range value. Also shown: the sampling distribution of the range parameter for heterogeneously fertilized plots that were either (D) Unfenced, or (H) Fenced. The distribution was calculated with a bootstrapping approach with maximum likelihood optimization.

