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7	Plant community responses to long-term fertilization: Changes in functional group
8	abundance drive changes in species richness
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20	experiment: KLG. Analyzed the data: TLD. Wrote the paper: TLD, KLG.

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### ABSTRACT

22 Declines in species richness due to fertilization are typically rapid and associated with increases in aboveground production. However, in a long-term experiment examining the 23 24 impacts of fertilization in an early successional community, we found it took 14 years for plant species richness to significantly decline in fertilized plots, despite fertilization causing a rapid 25 increase in aboveground production. To determine what accounted for this lag in the species 26 richness response, we examined several potential mechanisms. We found evidence suggesting 27 the abundance of one functional group—tall species with long-distance (runner) clonality— 28 drove changes in species richness, and we found little support for other mechanisms. Tall runner 29 species initially increased in abundance due to fertilization, then declined dramatically and were 30 not abundant again until later in the experiment when species richness and the combined biomass 31 32 of all other functional groups (non-tall runner) declined. Over 86% of the species found throughout the course of our study are non-tall runner, and there is a strong negative relationship 33 between non-tall runner and tall runner biomass throughout our study. We therefore suggest that 34 declines in species richness in the fertilized treatment are due to high tall runner abundance that 35 decreases the abundance and richness of non-tall runner species. By identifying the functional 36 group that drives declines in richness due to fertilization, our results help to elucidate how 37 fertilization decreases plant richness and also suggests that declines in richness due to 38 fertilization can be lessened by controlling the abundance of species with a tall runner growth 39 form. 40

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## **INTRODUCTION**

Increased fertilization via nitrogen deposition and other inputs is one of the most 42 important factors driving declines in species diversity in terrestrial communities (Clark and 43 Tilman 2008; Sala et al. 2000; Vitousek et al. 1997). A large number of fertilization 44 experiments, particularly in grasslands, show that fertilization decreases plant species richness 45 (DiTommaso and Aarssen 1989; Gough et al. 2000; Suding et al. 2005), especially when 46 fertilization leads to large increases in aboveground biomass production (Suding et al. 2005). 47 Many studies have examined the mechanisms by which fertilization decreases species richness 48 (Dickson and Foster 2011; Eilts et al. 2011), but only a few studies have examined whether 49 particular plant functional groups respond differently to fertilization and drive declines in species 50 richness (Clark et al. 2007; Gough et al. 2012; Suding et al. 2005). One cross-site study that has 51 52 examined the response of different herbaceous plant functional groups to fertilization finds that fertilization generally increases the abundance of one plant functional group—tall species with 53 long distance (runner) clonality (Gough et al. 2012). Motivated by results reported in Gough et 54 al. (2012), we examined how fertilization affects the abundance of different functional groups in 55 a long-term fertilization experiment. Specifically, we investigated whether increases in tall 56 runner abundance in response to fertilization could explain declines in species richness and the 57 abundance of other functional groups over time, and we also examined a number of other 58 alternate hypotheses. 59

If increases in the abundance of one functional group result in decreases in the
abundance of other functional groups, this should lead to the declines in species evenness often
observed after fertilization (Collins et al. 2008; Kempton 1979; Turkington et al. 2002). A metaanalysis by Hillebrand et al. (2007) found that fertilization can have no effect or even lead to

slight increases in species richness if no species dominate the community (high species
evenness). Hillebrand et al. (2007) also point out that herbivory can reduce the negative effects
of fertilization on species richness if herbivory decreases plant dominance. We sought to go
beyond an examination of how fertilization impacts community-level relative abundance and
explore how changes in the abundance of particular functional groups drive changes in diversity
following fertilization.

Here we present results from the first 23-years of an ongoing nitrogen addition 70 experiment conducted in a set of replicate successional old-field plant communities at the Long 71 72 Term Ecological Research project at the W.K. Kellogg Biological Station in southwestern Michigan. Fertilization did not significantly impact species richness until 14 years into the 73 experiment (Gough et al. 2000; Huberty et al. 1998), despite a significant increase in above-74 75 ground production. This lag in effect of fertilization on species richness provided an opportunity to examine how fertilization interacts with changes in species composition and functional group 76 abundance to affect species richness. 77

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#### **MATERIALS AND METHODS**

This experiment was conducted at Michigan State University's W.K. Kellogg Biological
Station (KBS), at the Long Term Ecological Research (LTER) site in agricultural ecology in
southwestern Michigan, USA (42°24' N latitude, 85°22' W longitude). Soils at this site are
Typic Hapludalfs, either fine-loamy, mixed, mesic Kalamazoo series or coarse-loamy, mixed,
mesic Oshtemo series (Mokma and Doolittle 1993). Temperatures range from an average
monthly high of 30°C in July to an average monthly low of -8°C in January. Precipitation (≈

1000mm annually) is spread fairly evenly throughout the year with somewhat less precipitationfalling in the winter than other seasons.

The experiment was established in 1989 in six replicate plots (each 105 m x 87 m) of the 88 successional treatment of the KBS LTER. Replicate fields were plowed in the spring following 89 20 + years of continuous corn production, and natural succession was then allowed to occur with 90 occasional prescribed burns (see below). Succession led to herbaceous vegetation dominated by 91 Solidago (goldenrod) species (S. altissima and S. graminifolia), and to a lesser extent by 92 Apocynum cannabinum (dogbane), and Elymus repens (quackgrass)—all tall, highly clonal 93 species (Huberty et al. 1998). Two 5 m x 5 m subplots were established in the north end of each 94 plot and one of the subplots was fertilized with 12 g N m<sup>-2</sup> in early summer (typically mid-June), 95 corresponding to the timing of fertilizer application in the surrounding row-crop treatment of the 96 97 LTER. Fertilizer was added as ammonium nitrate pellets. Information about soil nitrogen levels throughout the experiment can be found in Fig. S1. Further information about the site and the 98 experiment can be found in Huberty et al. (1998) and Grman et al. (2010). 99 100 Plant species composition and biomass was determined annually from both the fertilized and unfertilized (control) subplots. To avoid edge effects, harvested quadrats were located in the 101 central (3 m x 3 m area) of the subplot; the location of the harvested quadrat was shifted each 102 year so that harvested areas were not resampled for 5 years. Harvests were completed at peak 103 biomass (typically early August) and all vegetation rooted in the harvested quadrat was clipped 104 105 at ground level. Harvested biomass was separated into live and dead biomass; live biomass was separated by species, dried at  $60^{\circ}$ C to constant weight, and weighed with 0.01 g accuracy. 106 Species richness and abundance were calculated from the harvested live biomass. In the first 107 three years of the study (1989-1991), the area of the harvested quadrat varied (0.2  $m^2$  in 1989 and 108

0.3 m<sup>2</sup> in 1990-1991), but since 1992 the quadrat has been standardized at 1 m<sup>2</sup> (0.5 m x 2 m).
We show the effects of fertilization on species richness for all years (Fig. 1), but only use data
from 1992 onwards for all other analyses.

We classified species into three height groups based on their position in the canopy (top 112 third = tall, middle third = mid, bottom third = short) and three clonal growth form groups 113 (runners = species with a spreading morphology that can send up new shoots some distance from 114 the original clone; clumpers = species which produce shoots adjacent to the original clone, non-115 clonal = species that do not spread) by utilizing height and clonality information in McGregor et 116 al. (1986) and through personal observations of species' height (a species could have the 117 potential to grow taller at some locations in its range rather than at our site). This classification 118 is similar to that used by Gough et al. (2012) and Cleland et al. (2008), except that we have 119 120 differentiated between mid and tall canopy species in our analysis.

Site Management: Woody species began to invade the site a few years after the start of 121 the experiment (Foster and Gross 1999), and by 1997 trees like *Robinia pseudoacacia* (black 122 123 locust) and *Rhus typhina* (stag horn sumac) became common in some replicates. Because the intent of this successional treatment was to provide an herbaceous-dominated reference 124 community for comparison with the crop treatments on the KBS LTER, management was 125 initiated to remove woody invaders from these plots and prevent reinvasion. This included 126 initiation of whole-plot annual prescribed burning in 1997 (spring) to control further colonization 127 by woody species and cutting and removing woody stems. Details of the management of these 128 fields can be found at (http://lter.kbs.msu.edu/). We did not find that species richness or the 129 effect of fertilization on species richness was significantly affected by burning (Fig. S2). 130

131	Statistical Analysis: All statistical analyses were completed using PROC MIXED in
132	SAS 9.2 (Cary, NC, USA). We sought to determine how the effects of fertilization on plant
133	species richness and the biomass of different functional groups varied through time, and whether
134	the biomass of tall runner and non-tall runner species could help to explain the changes in plant
135	species richness over time. Therefore, we performed split-plot, repeated-measure Analysis of
136	Variance (RM-ANOVA) for analyses, except where a covariate such as tall runner biomass or
137	non-tall runner biomass was included in which case we performed split-plot, repeated-measure
138	Analysis of Covariance (RM-ANCOVA). We used a Kenward-Rogers correction in all analyses,
139	which led to reduced degrees of freedom and to decimals in the degrees of freedom due to
140	missing data (biomass was not collected from one quadrat in 1997).
141	We tested the data for normality and transformed the data if necessary to improve
142	normality (data are transformed in figures in these cases). We treated each plot as a separate
143	block. The blocking term was added as a random effect to all analyses to remove any effects of
144	spatial variation and is not reported.
145	
146	RESULTS
147	Over the first 13 years of the experiment (1989-2001), fertilization did not significantly
148	affect plant species richness (Fig. 1), despite the fact that fertilization significantly increased total
149	biomass by 64% during this period (Fig. 2A). In 2002, 14 years after the fertilization treatment
150	was initiated, species richness significantly declined in the fertilized plots and remained lower
151	most years thereafter (Fig. 1; year x fertilizer interaction). We set out to determine why
152	fertilization did not decrease species richness until 2002, even though total biomass was quickly
153	increased by fertilization.

154 The observed decline in species richness in fertilized plots occurred when fertilization led to a sustained increase in the abundance of species with a tall runner growth form. Tall runner 155 biomass initially increased due to fertilization, then declined sharply in 1995, and then showed a 156 sustained increase at the same time species richness declined (Fig. 2B). Two species in the 157 genus Solidago (S. altissima and S. graminifolia) made up most of the tall runner biomass in the 158 fertilized treatment in the first part of the experiment (1989-2001). In the latter part of the 159 experiment (2002-2011) when tall-runners again dominated the fertilized plots, Solidago species 160 were again abundant, but made up less of the tall runner biomass (Fig. 2B). The biomass of all 161 162 other functional groups (non-tall runners) was not significantly affected by fertilization until 2005, but was significantly decreased by fertilization in most years from 2005 onwards (Fig. 163 2C). The timing of the decline in biomass of these other functional groups generally 164 165 corresponded to when we observed a significant decline in species richness (Fig.1). The change in the relative abundance of tall runners in response to fertilization can also be clearly seen when 166 examining the percent of total biomass made up of tall runner species over the course of the 167 168 experiment (Fig 2D). No functional group other than tall runners increased in biomass in response to fertilization (Fig. S3). 169

It appears fertilization decreased species richness when tall runner biomass remained high for long enough to lead to decreases in non-tall runner biomass. Specifically, there was a significant negative relationship between tall runner biomass and non-tall runner biomass (Fig 3A). Over 86% of species observed from 1992 onwards had a non-tall runner growth form, which helps to explain why species richness declined when tall runner biomass increased—and non-tall runner biomass decreased. There was a significant positive relationship between nontall runner biomass and species richness (Fig. 3B), and a significant negative relationship

between tall runner biomass and species richness (Fig. 3C; note that the fertilization term in the
model is not significant when tall runner biomass is included as a covariate). The biomass of no
other functional group or species accounted for these observed changes in species richness (Fig.
S3-S4). Also, the relationship between total biomass and species richness was only weakly
significant (Fig. 3D), suggesting that changes in the biomass of the tall runner and non-tall
runner functional groups were driving changes in richness rather than changes in total biomass.

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#### DISCUSSION

185 While declines in species richness are commonly observed in response to fertilization in grasslands (Clark et al. 2007; Suding et al. 2005), typically these responses occur quickly and are 186 correlated with increases in community biomass (Suding et al. 2005). We are not aware of any 187 188 other study that has documented such a long-term temporal lag (14 years) in the species richness response to fertilization when there is a strong fertilization effect on above-ground production. 189 The magnitude of the total biomass response to fertilization does not appear to provide a 190 191 sufficient explanation as to why it took 14 years for species richness to decline in this experiment. Specifically, total biomass in the fertilized treatment was significantly higher than 192 the unfertilized treatment in most years of the experiment, except drought years, as suggested by 193 the significant effect of fertilization and the non-significant year x fertilizer interaction (Fig. 2A). 194 Also, the effects of fertilizer on total biomass in our study were similar in magnitude to other 195 196 fertilization experiments (Gough et al. 2012).

Instead, we found that increases in the biomass of a particular functional group—species with a tall runner growth form—rather than increases in total biomass, appeared to drive declines in species richness by decreasing the average biomass of the 86% of species that are not tall

200 runner. Gough et al. (2012) also found that the tall runner functional group consistently 201 increased in abundance in response to fertilization across 37 experiments in North American grasslands. Interestingly, Gough et al. (2012) found a consistent response of the tall runner 202 203 functional group across many locations, but the response was generally driven by a single—but different—species at each location. The response at our site was primarily driven by *Solidago* 204 species, primarily S. altissima. Solidago species made up much of the tall runner biomass, and 205 provided almost as good a fit to species richness as did tall runner biomass (Fig. S5). However, 206 we focus on tall runner species because they provide the strongest fit with species richness (Fig. 207 208 S5) and because of the generality of their response to fertilization across sites (Gough et al. 2012). 209

The Gough et al. (2012) study did not find a significant relationship between species 210 211 richness and the response of tall runner species to fertilization, but they noted that the relative abundance ratios used in their analyses may have masked such a relationship. In addition, the 212 Gough et al. (2012) meta-analysis used data from only one year of each of the 37 experiments 213 214 (last year of data available from each site in Cleland et al. 2008 data set). This would also have masked the relationship between tall runner species and richness (ex. used the 2001 data from 215 our study, in which fertilization did not significantly decrease richness). Our results thus expand 216 upon Gough et al. (2012) by examining the effects of tall runner species through time and by 217 using analyses that reveal how changes in functional group abundance over time can drive 218 219 community responses to fertilization.

The observation that tall runner biomass appeared to drive changes in species richness raises two obvious, but difficult to answer questions: 1) Why did the initial spike in tall runner biomass in the first part of the experiment not cause a decrease in species richness?; and 2) What

223 caused the precipitous decline in abundance of tall runner species in the fertilized treatment in 224 1995? The answer to the first question may be that high tall runner biomass was not sustained for long enough in the first part of the experiment to decrease non-tall runner biomass and 225 226 species richness. Transient dynamics can occur whereby inferior competitors persist until resource levels equilibrate (Tilman 1988) or disturbance ceases (Huston 1979). We suggest that 227 one year of high tall runner biomass in 1994 was not enough to exclude many species—figure 228 2C shows that non-tall runner biomass decreased in 1994 as tall runner biomass reached its 229 maximum, but then recovered in 1995 when tall runner biomass declined. 230

231 The answer to the second question requires a bit more speculation and inference from other experiments that have examined the impacts of insect herbivores on plant communities and 232 how this varies in response to fertilization. Most of the tall-runner biomass at the beginning of 233 234 the experiment was composed of *Solidago* species; in the fertilized treatment S. altissima and S. graminifolia made up over 85% of the tall-runner biomass and 65% of the total biomass from 235 1992-1994. These are both native species that are vulnerable to predation by insect herbivores 236 237 (Carson and Root 1999), and we suggest a Solidago-specific insect herbivore may have decreased *Solidago* biomass in 1995. Work by Carson and Root (2000) provides a good example 238 of how this could occur-in their study an outbreak of a Solidago-specific insect herbivore 239 (Microrhopala vittata) strongly reduced Solidago abundance for several years relative to insect 240 exclusion treatments and led to increases in plant species richness. Other studies have shown 241 that a variety of insect herbivores (including insects in the *Trirhabda* genus) can strongly reduce 242 Solidago abundance (Carson and Root 1999; Maddox and Root 1990). Although we have no 243 direct evidence of a1995 insect outbreak, Trirhabda species were observed in the plots during 244 245 this time (Katherine Gross personal observations). It has also been shown that insects often

preferentially feed on fertilized vegetation (Uriarte 2000), which may help to explain why the decline in *Solidago* biomass was so large in the fertilized treatment. More broadly, grazing by small mammals has also been shown to reduce the negative effects of fertilization on plant species richness (Gough and Grace 1998; Inouye et al. 1987). Regardless of what accounts for the rapid decline and delayed re-establishment of tall runner species in this system, our results suggest that changes in tall runner abundance related to fertilization are an important driver of declines in plant species richness following fertilization.

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## 254 Other potential hypotheses

We explored whether other hypotheses beyond the abundance of tall runner species could 255 explain both the magnitude and the delay of the species richness response in our study. 256 257 However, we could not find support for any other explanations. Fire has been shown to be an important driver of plant biomass and species richness in tall-grass prairie (Collins et al. 1998). 258 In our experiment, fire was introduced in 1997 to control woody species invasion, and while we 259 260 do not have unburned control plots, we find no evidence that fire drove changes in species richness (Fig. S2). Similarly, increased litter biomass in response to fertilization has been shown 261 to be an important limit on species richness in grasslands (Foster and Gross 1998). We examined 262 whether litter biomass was correlated with species richness in our study, but found no significant 263 relationship (Fig. S6). We also examined whether the abundance of other functional groups (ex. 264 legumes, grasses) could explain changes in species richness over time, but found little support 265 for other functional groups driving changes in richness over time (Fig. S4). Lastly, fertilization 266 has been shown to alter colonization rates (Tilman 1993), which could be important in our 267

268	successional fields (Huberty et al. 1998). However, fertilization had little effect on species
269	colonization rates (gain or loss of species) in our study (Fig. S7).

While other factors may have contributed to the lag in effects of fertilization on species richness, it seems that the consistent increase in tall runner biomass in the latter part of the experiment and the associated decrease in non-tall runner biomass is the most parsimonious explanation for this lag. However, as with every time series experiments, we cannot rule out the possibility of unmeasured variables affecting plant species richness over time and future experiments should be completed to isolate the effects of tall runner biomass on species richness.

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## 277 General relevance

The results of our study provide insights to understanding the responses of plant 278 279 communities to long term perturbations. We have shown that particular functional groups can play an important role in controlling plant species richness under fertilized conditions, and we 280 discuss below how this may be important when managing sites with high soil fertility for high 281 282 plant diversity. While other studies have shown that species dominance can be a driver of diversity (Hillebrand et al. 2007), the role of particular functional groups in controlling richness 283 under fertilized conditions is less well understood. For example, Clark et al. (2007) presented 284 evidence that sites with higher C<sub>4</sub> graminoid abundance showed strong responses to fertilization 285 and this correlated with greater species diversity loss. Although the abundance of  $C_4$  graminoid 286 species was initially high in our study, this functional group rapidly declined in abundance (Fig. 287 S4), and the abundance of this functional group did not correspond to changes in species richness 288 in response to fertilization. Suding et al. (2005) presented evidence that greater shifts in 289 290 abundance of functional groups was correlated with greater species losses, and they also did not

find C<sub>4</sub> graminoids were important in determining species losses. Instead, they found that legumes were the most likely functional group to be lost due to fertilization. We also found that legume species were lost in the fertilized treatment, but this only explained a small amount of total species losses due to fertilization. We suggest that height and clonality functional groups provide greater predictive power for the effects of fertilization on plant species richness than do growth form (graminoid, non-leguminious forb, leguminous forb) or photosynthetic pathway (C<sub>4</sub> and C<sub>3</sub>) functional groups.

Few studies have considered how to prevent species losses under fertilized conditions 298 (but see Collins et al. 1998, Foster et al. 2010, Symstad and Jonas 2011). The studies that have 299 considered how to prevent losses in species richness have generally focused on the role of 300 disturbance in preventing losses. Disturbances such as grazing or having have been shown to 301 302 increase plant diversity under high productivity conditions (Collins et al. 1998; Foster et al. 2010; Hillebrand et al. 2007; Symstad and Jonas 2011), but these studies give little detail on the 303 mechanisms by which disturbances increase plant diversity. Understanding these mechanisms 304 305 will become increasingly important if grasslands are increasingly fertilized and disturbed as part of cellulosic biofuel production (Gelfand et al. 2013). 306

Disturbances may help to prevent losses of species due to fertilization because fertilization increases the abundance of dominant species (Turkington et al. 2002; Collins et al. 2008; Fig. S8), and disturbance may decrease the abundance of dominant species (Hillebrand et al. 2007). A meta-analysis across many studies shows that fertilization only decreases species richness when a few species are highly abundant and evenness is low (Hillebrand et al. 2007). These results are similar to our finding of a negative relationship between tall runner and non-tall runner biomass and of plant species richness not declining until sustained increases occur in tall

314 runner biomass. There are several studies that suggest removing dominant (Gurevitch and Unnasch 1989) or clonal species (Eilts et al. 2011; Reynolds et al. 2007) generally removes the 315 negative effects of fertilization on species richness. In contrast, Lepš (1999) found that removal 316 317 of dominant species does not necessarily prevent a decrease in species richness after fertilization, although other species increased in abundance after dominant removal in this study and may 318 have filled the role of dominant competitors. If fertilization is primarily acting by increasing the 319 320 abundance of dominant species which thereby reduces the abundance of subdominant species, then future research should examine how changes in the abundance of dominant and 321 322 subdominant species affect species richness rather than how changes in total biomass affect species richness. 323

The mechanisms by which highly abundant species decrease plant richness under 324 325 fertilized conditions also need to be determined. One might expect that increases in the abundance of dominant species would decrease light levels and thereby lead to decreases in the 326 abundance of non-tall runner species. Light manipulation experiments in the presence and 327 328 absence of tall runner species would be necessary to fully address the role of light competition. However, we suggest that all tall-statured species—not just tall runner species—would increase 329 in abundance under fertilized conditions if light competition was the only factor causing changes 330 in functional group biomass. We found fertilization decreased or had no effect on the biomass of 331 tall-statured species without the runner growth form (Fig. S3), suggesting that plants competed 332 for additional resources beyond light (see also Dickson and Foster 2011). 333

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335 Conclusions

336	Overall, our results suggest that fertilization decreases plant richness by increasing the
337	abundance of tall runner species and thereby decreasing the abundance of other functional
338	groups. Other than Gough et al. (2012), the effects of fertilization on tall runner biomass has not
339	been examined, and future studies should examine the role of tall runner species in driving
340	declines in plant richness in other habitats.
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<ul><li>342</li><li>343</li><li>344</li><li>345</li><li>346</li></ul>	ACKNOWLEDGEMENTS We gratefully acknowledge Carol Baker, Pam Moseley, and many field workers whom have maintained and sampled the experiment over its long history. We received helpful comments on earlier drafts of this manuscript from Bryan Foster, Laura Gough, and two anonymous reviewers. The work was supported by several National Science Foundation grants: DEB 87-0232 and

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**FIGURE LEGENDS** 432 Figure 1. Effects of fertilization and year on plant species richness (species richness refers to the 433 number of species per sample area; sample area was less than  $1 \text{ m}^2$  in first three years and  $1 \text{ m}^2$ 434 435 from 1992 onwards, see Methods for details; values are means + 1 SE; N=6). 436 Figure 2. Effects of fertilization and year on (A) total community biomass, (B) biomass of tall 437 438 runner species and Solidago species, (C) biomass of all species other than tall runner species, and (D) percentage of total biomass made up of tall runner species (values are mean + 1 SE; N=6). 439 440 Figure 3. The relationship between (A) tall runner biomass and the biomass of all other species, 441 (B) the biomass of non-tall runner species and plant species richness, (C) tall runner biomass and 442

443 plant species richness, and (D) total biomass and plant species richness (note that non-tall runner

biomass is on a square root scale).



Figure 1.



Figure 2.



Figure 3.