

Livestock grazing regulates ecosystem multifunctionality in semiarid grassland

Journal:	<i>Functional Ecology</i>
Manuscript ID	FE-2018-00771
Manuscript Type:	Research Article
Date Submitted by the Author:	08-Aug-2018
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Livestock grazing regulates ecosystem multifunctionality in semiarid grassland

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Running headline: What drives grassland ecosystem multifunctionality?

Abstract

- 1 1. Livestock grazing has been shown to alter the structure and functions of grassland
2 ecosystems. It is well acknowledged that grazing pressure is one of the strongest drivers
3 of ecosystem-level effects of grazing, but few studies have assessed how grazing pressure
4 impacts grassland biodiversity and ecosystem multifunctionality (EMF).
- 5 2. Here, we assessed how different metrics of biodiversity (i.e. plants and soil microbes) and
6 EMF responded to seven different grazing treatments based on an 11-year field
7 experiment in semi-arid Inner Mongolian steppe.
- 8 3. We found that soil organic carbon, plant-available nitrogen, and plant functional diversity
9 all decreased even at low grazing pressure, while aboveground primary production and
10 bacterial abundance decreased only at high levels of grazing pressure.
- 11 4. Structural equation models revealed that EMF was driven by direct effects of grazing,
12 rather than the effects of grazing on plant or microbial community composition. Grazing
13 effects on plant functional diversity and soil microbial abundance did have moderate
14 effects on EMF, while plant richness did not.
- 15 5. *Synthesis*. Our results showed ecosystem functions differ in their sensitivity to grazing
16 pressure, requiring a low grazing threshold to achieve multiple goals in the Eurasian
17 steppe.

18

19 **Key-words:** species richness, functional diversity, soil microbes, threshold, grazing pressure,
20 semi-arid grassland

21 Across the planet, grasslands are the most common land cover type. These ecosystems
22 support over 2.5 billion people, most of whom directly rely on ecosystem services for
23 survival and livelihood (MEA 2005; Reynolds *et al.* 2007; Briske 2017; Evans *et al.* 2017).
24 However, grasslands are one of the most vulnerable ecosystems, facing degradation of plant
25 diversity, soils, and ecosystem services (MEA 2005; Teague & Barnes 2017). Development of
26 sustainable grazing systems that promote ecosystem resilience, enhance or maintain plant
27 diversity, increase soil health, and maintain ecosystem multifunctionality (EMF) and delivery
28 of multiple ecosystem services is a global concern (MEA 2005; Sala *et al.* 2017; Teague &
29 Barnes 2017). Balancing these multiple objectives can be challenging because tradeoffs are
30 common across multiple grassland management goals (MEA 2005; Briske *et al.* 2011;
31 Maestre *et al.* 2012; Jing *et al.* 2015). For example, many ecosystem services are linked with
32 plant diversity, yet management for other ecosystem services may reduce plant diversity (e.g.
33 maximizing productivity by promoting dominant plant species) (Bullock *et al.* 2011). Adding
34 to this complexity, ecological impacts of grazing can be highly variable, depending on
35 interactions between grazing management practices and environmental conditions (Briske *et*
36 *al.* 2011). This complexity makes it difficult to set prescriptions for livestock grazing
37 practices.

38 Of all aspects of grazing practices, livestock stocking density has the strongest
39 ecosystem-level impacts (Briske *et al.* 2011). However, best management practices tend to
40 focus on a subset of ecosystem characteristics. For example, moderate grazing pressure
41 maximizes plant productivity in semi-arid Eurasian steppe grasslands (Liu *et al.* 2015; Li *et*
42 *al.* 2017), and improves plant diversity in relatively productive grasslands, but reduces plant

43 diversity in less productive grasslands (Huston 1979; Kondoh & Williams 2001).
44 Understanding the effects of herbivore density, or grazing pressure, on the ability of an
45 ecosystem to deliver multiple functions (hereafter Ecosystem Multifunctionality, EMF) is
46 critical to determine sustainable grazing practices and the delivery of multiple ecosystem
47 services (Schonbach *et al.* 2011; Stein *et al.* 2016).

48 Approximately 45% of variation in EMF is explained through combined effects of
49 above-and belowground biodiversity (Jing *et al.* 2015). Thus, the development of best
50 management practices requires improving our understanding of the influence of herbivory on
51 plant and microbial communities, and their effects on multiple ecosystem processes (Bardgett
52 & Wardle 2003; Harrison & Bardgett 2010; Sitters & Venterink 2015; Evans *et al.* 2017; Liu
53 *et al.* 2018). Herbivory can strongly affect plant community structure and function (Diaz *et al.*
54 2007b; Stein *et al.* 2016), through the direct effects of herbivores on plants, and
55 grazing-induced changes in soil nutrients and fungal communities (Chen *et al.* 2017; Ren *et*
56 *al.* 2018). Lack of grazing can decrease species diversity because of competitive exclusion
57 and light limitation (Borer *et al.* 2014). These changes in plant community composition can
58 lead to large shifts in soil microbial communities and processes (Stein *et al.* 2016; Wilson *et*
59 *al.* 2018). Grazing-induced changes in plant functional traits can be particularly important in
60 understanding ecosystem multifunctionality. High grazing pressure has been shown to reduce
61 functional diversity (FD) (Gross *et al.* 2007; Gross *et al.* 2014; Baert *et al.* 2016; Li *et al.*
62 2017). Relatively high FD may benefit an ecosystem by enhancing plant community
63 complementarity in resource acquisition and utilization and promoting community resilience
64 and resistance. Diverse plant communities and community FD are strongly related to multiple

65 ecosystem functions (Petchey & Gaston 2002; Forrestel *et al.* 2017), and high FD can
66 maintain high EMF and ecosystem resilience (Valencia *et al.* 2015).

67 Soil biota are direct mediators of carbon, nitrogen, and phosphorus cycles, and are
68 therefore important drivers of plant diversity and ecosystem productivity (van der Heijden,
69 2008; Wurzburger & Brookshire 2017). Soil biota are strongly affected by herbivore grazing
70 (Barto & Rillig 2010; Chen *et al.* 2013; Liu *et al.* 2015; Eldridge *et al.* 2017) through
71 multiple pathways, including changes in plant community composition, soil nutrients,
72 moisture, and compaction. In addition, carbon allocation to roots and root exudates, directly
73 alter the abundance of arbuscular mycorrhizal (AM) fungi and other soil organisms (van der
74 Heyde *et al.* 2017; Chen *et al.* 2017; Ren *et al.* 2018; Wilson *et al.* 2018).

75 While there is still considerable debate on the ecosystem-level effects of specific grazing
76 practices (e.g. rotational vs. continuous grazing), it is well documented that livestock stocking
77 density (grazing pressure) has strong impacts on all aspects of the ecosystem. However, the
78 ideal grazing pressure for any given system is largely unresolved (Briske 2017).
79 Understanding the effects of grazing pressure on EMF is critical to determine sustainable
80 grazing practices (Schonbach *et al.* 2011; Stein *et al.* 2016). Here, we examined how plant
81 species richness, plant FD (including five functional traits: plant species height, specific leaf
82 area, leaf dry matter content, leaf nitrogen content, and stem:leaf ratio), soil microbes,
83 grazing pressure, and soil factors (soil moisture and pH) influenced EMF. In our study, we
84 utilized EMF to summarize five key ecosystem functions and related variables: (1)
85 aboveground biomass, (2) plant nitrogen (nitrogen pools in aboveground biomass), (3)
86 plant-available nitrogen, (4) plant-available phosphorus, and (5) soil organic carbon. Our

87 experiment investigated the following: (i) the effects of grazing pressure on EMF (e.g., does
88 moderate grazing pressure maintain or improve EMF, according to the intermediate
89 disturbance hypothesis? (Hanke *et al.* 2014)); and (ii) the extent that grazing directly alters
90 EMF, versus indirectly affects EMF through changes in plant and microbial communities

91 **Methods**

92 **Site description**

93 Our study area is located in Inner Mongolia steppe (Bai *et al.* 2004), ranging in elevation
94 from 1200 to 1280m, with a mean annual precipitation of 346.1mm falling mainly in the
95 growing season from May to September, and with a mean annual temperature of 0.3°C, with
96 the lowest mean monthly temperatures ranging from -21.6 °C in January to the highest 19.0
97 °C in July. The study area has a history of long-term grazing at moderate to heavy grazing
98 pressure, but livestock were excluded from this area two years prior to the start of the
99 experiment in 2005. Each year, sheep are in the field from June to September (~95 days), in
100 accordance with the local summer grazing season. Soil is classified as Calcic Chernozem
101 (IUSS Working Group WRB 2006). Approximately 36 vascular plant species typically occur
102 in these grasslands (8 of them are very rare), grouped by functional characteristics: perennial
103 rhizomatous grasses, perennial bunchgrasses, perennial forbs, and annual/biennial grasses
104 (Sasaki *et al.* 2009; Wu *et al.* 2015). The dominant perennial rhizomatous grass *Leymus*
105 *chinensis* and the perennial bunchgrass *Stipa grandis*, together account for approximately
106 75% of total aboveground biomass production (Li *et al.* 2017).

107 **Grazing treatments**

108 Our project was designed to assess the impacts of grazing at temporal and spatial scales
109 that are both relevant to management, and that can capture ecosystem- and landscape-scale
110 effects of grazing. There has been strong support for this approach, emphasizing that
111 small-scale plots: (1) often yield different results than ecosystem-scale plots, (2) do not
112 address heterogeneity in grazing/disturbance/management across the landscape, and (3) are
113 impossible to scale-up to inform management decisions (Carpenter 1996, Schindler 1998,
114 Schmitz 2005, Osenberg et al. 2006, Fraterrigo and Rusak 2008). These ecosystem-scale
115 studies require large land areas and high-levels of logistics, tend to be expensive, and thus
116 there is limited ability to replicate large-scale experimental plots. In fact, reviews of such
117 large-scale experiments suggest that because of the difficulty of replication, if multiple
118 large-scale plots are feasible, it is more valuable to include additional treatments, rather than
119 replicating the same treatment (Schindler 1998). Strong statistical inferences can be drawn by
120 focusing on a regression-based experimental design, in which multiple levels of a treatment
121 are applied (with or without replication). This regression approach is more powerful
122 statistically than replicated ANOVA-based designs, and allows for research that is more
123 relevant to both management and predictive ecology, by assessing how the effects of the
124 treatment vary with level of the treatment (Cottingham et al. 2005). This regression approach
125 is particularly effective for a broad array of management-scale questions, ranging from
126 effects of grazing, to effects of precipitation change (Brandsby et al 1998, Beier et al. 2012).

127 Following a regression-based design, in April 2005, a grazing experiment covering 160
128 ha was established and maintained for 11 years (Schonbach *et al.* 2011). The grazing
129 manipulations occurred at two site types (flat or sloped), with each site type containing 7

130 plots that were randomly assigned to 7 grazing pressures (GP) (GP = 0, 1.5, 3.0, 4.5, 6.0,
131 7.5, or 9.0 sheep ha⁻¹). These two site types have similar response to grazing treatments
132 (Supplementary Fig. 5 a, b), and thus were pooled in the statistical analyses.

133 Our study utilized non-lactating female sheep with an average live weight of 35 kg. The
134 plots were ~2 ha in size, except for the lowest grazing pressure (1.5 sheep ha⁻¹), which was
135 ~4 ha to ensure at least six sheep per plot. There was no significant difference between plots
136 in either plant species composition or relative abundance of plant species before initiation of
137 our study, but species composition and community structure did change in response to
138 grazing treatments (see Li *et al.* 2017).

139 **Plant and soil sampling**

140 All plant and soil measures were collected at the end of the 2015 growing season, a year
141 with higher annual precipitation and temperature than average. Higher precipitation is likely
142 linked with higher species diversity. Samples were collected from nine randomly placed 1m²
143 quadrats within each treatment plot.

144 *Plant sampling*

145 In these plots, we assessed plant species composition (% cover) and richness (number of
146 plant species) . Table 1 contains a list of all vascular plant species identified. To measure
147 biomass throughout the growing season, we established three exclosure cages (2 × 3m) in
148 each plot before sheep began grazing. From June through September, aboveground biomass
149 was clipped in a 1m² quadrat from both inside and outside of each exclosure. After each
150 monthly clipping, exclosures were moved. Annual aboveground net primary productivity

151 (ANPP) inside (i) and outside (o) exclusures in grazed plots was calculated with the formula:
152 $ANPP = W_{1o} + (W_{2i} - W_{1o}) + (W_{3i} - W_{2o}) + (W_{4i} - W_{3o})$. Where W_i represents standing
153 plant biomass at the start of each month (1 = June, 2 = July, 3 = August, and 4 = September).
154 The biomass is presented on a dry weight basis. We determined plant-tissue N concentration
155 using the Kjeldahl method (Kjeltec 8100 Analyser Unit, FOSS, Sweden).

156 *Soil sampling*

157 Soil samples (diameter 3 cm, depth 10 cm) were collected at the end of the 2015 growing
158 season from nine randomly placed locations in each plot. Subsamples for soil organic carbon
159 and plant-available phosphorus and nitrogen analyses were air-dried, sieved through a 2mm
160 mesh and ground to a fine powder. Subsamples were also separated for soil moisture, soil pH,
161 and soil microbial analyses (AM extra-radical hyphae, saprophytic fungi, and bacteria). Soil
162 bulk density at 0-10 cm depth was measured using a cutting ring (volume of 100 cm³).

163 **Soil properties determination**

164 Plant-available P was measured by the Olsen method. Soil organic C was analyzed by the
165 dry combustion method (Multi N/C 2100, Analytik jena, Germany). Plant-available N was
166 also measured by Multi N/C 2100, from extractions with 50 mL of 2 mol L⁻¹ K₂SO₄ from 10
167 g fresh field soil. To determine soil moisture content, twenty grams of fresh soil was weighed
168 before and after oven-drying at 105 °C for 24 h. Ten grams of field soil was mixed with 25
169 mL of 1 mol L⁻¹ KCl solution to measure pH using a pH meter (PB-10, Sartorius, Germany).
170 Extra-radical hyphal length densities of AM fungi were extracted from soil using the
171 membrane filter technique and the gridline intercept method under a microscope at 200X

172 magnification (Jakobsen *et al.* 1992). The biomass of soil bacteria and saprophytic fungi were
173 calculated using phospholipid fatty acid (PLFA) analysis. Qualitative and quantitative fatty
174 acid analyses were performed using an Agilent 6890 gas chromatograph (Agilent
175 Technologies, USA) and Sherlock software (MIDI, USA). The PLFA biomarkers a15:0,
176 i15:0, i16:0, 16:1x7, i17:0, a17:0, 17:0, cy17:0, cy19:0 were selected to represent soil
177 bacteria, and 18:2 ω 6c was selected to represent saprophytic fungi (Moore-Kucera & Dick
178 2008).

179 **Quantifying functional diversity**

180 To test grazing effects on plant functional traits, we coupled our species composition
181 data with quantitative values for species functional traits. Functional trait data were collected
182 from thirty plants per species, grown in non-grazed plots. We focused on five functional plant
183 trait responses: plant species height (SH), specific leaf area (SLA), leaf dry matter content
184 (LDMC), leaf nitrogen content (LNC), stem:leaf ratio. These traits were chosen because they
185 link to plant nutrient acquisition and utilization, belowground interactions with soil microbes
186 and fauna, and because they tend to be indicators of plant sensitivity to grazing. Plants with
187 low SLA and high LNC are negatively affected by intense grazing pressures (Garnier *et al.*
188 2004; Li *et al.* 2017). To allow for comparison across traits that vary in units and magnitudes,
189 we standardized plant trait values by transforming them with $\log_{10}(x + 1)$. We then averaged
190 values by species and used averages in calculations of functional diversity.

191 There are several ways to calculate plant functional diversity. While some, such as the
192 community weighted mean focus on single traits, we opted for Mason functional diversity

193 index, which is an integrated measure of all assessed plant functional traits at the
 194 community-level (Lavorel *et al.* 2008; Valencia *et al.* 2015). The Mason functional diversity
 195 index can represent overall community-level trait values by accounting for the abundance of
 196 each species in each plot (Mason *et al.* 2003; Mori *et al.* 2017).

197 $FD\alpha$ represents Mason functional diversity index:

$$198 \mathbf{FD}\alpha = \sum_{i=1}^S P_i (x_i - \bar{x}) \quad (1)$$

199 x_i represents the mean trait value of species i , $\bar{x} = \sum_{i=1}^S P_i x_i$ represents the mean trait value of
 200 whole plant community. P_i represents the relative abundance of species i in the whole plant
 201 community, and S represents the number of species in the whole community (Mason *et al.*
 202 2003, 2011, 2013; Mason *et al.* 2005).

203 **Quantifying Ecosystem Multifunctionality (EMF)**

204 EMF index is used as an integrated measure of a system's ability to sustain multiple
 205 functions simultaneously. Variables that we included in our calculation of EMF are: (1) plant
 206 aboveground biomass, (2) plant-tissue nitrogen content, (3) plant-available nitrogen, (4)
 207 plant-available phosphorus, and (5) soil organic carbon. All of these variables are crucial
 208 drivers of ecosystem functioning (Jing *et al.* 2015) (Delgado-Baquerizo *et al.* 2017a, b), as
 209 well as key factors for plant and soil health. Several methods can be used to calculate EMF,
 210 each with merits and faults. Here, we used a common method, “averaging approach (EMF
 211 index),” to calculate ecosystem multifunctionality (Hooper & Vitousek 1998; Maestre *et al.*
 212 2012). The “averaging approach (EMF index)” assesses the average effect of diversity across
 213 a suite of functions, with values of functions standardized. Because it averages, it cannot

214 distinguish between one function being provided at a high level and another being provided at
 215 a low level, vs. two functions being provided at an intermediate level (Byrnes *et al.* 2014).
 216 Thus, we have supplemented this averaging approach with a threshold analysis approach.
 217 Threshold analysis specifies how many functions are provided above 50% of the maximum
 218 provision. Together, these give a sense of the extent that diversity influences the average
 219 provisioning of ecosystem functions, and the number of functions provided at a high level.

220 To calculate EMF, we standardized EMF values ranging from 0 to 1
 221 ($f(x) = (x - \min(x)) / (\max(x) - \min(x))$), providing a unifying dimension across multiple
 222 functions (Gamfeldt & Roger 2017).

$$223 \text{EMF}^{\alpha} = (\sum_{i=1}^F g(r_i(f_i))) / F \quad (3)$$

224 EMF^{α} represents ecosystem multifunctionality index, f_i represents the value of function i ,
 225 r_i represents mathematical function for transforming the f_i value into a positive value, g
 226 represents the standardizing of all values, and F represents the number of measured functions.

227 **Statistical analysis**

228 All statistical analyses were performed using SAS Version 9.1 (SAS Institute Inc., Cary,
 229 NC, USA) and R version 3.3.1 (R Foundation for Statistical Computing, Vienna, Austria,
 230 2013). For all analyses, data were $\log_{10}(x+1)$ transformed to ensure normality and
 231 homogeneity, as confirmed by the Shapiro-wilk test. Two replicates per grazing level (slope
 232 vs. flat areas) were averaged and used in analyses. Ordinary least squares (OLS) regressions
 233 were used to assess how grazing pressure correlated with plant functional diversity and each

234 ecosystem function. Adjusted R^2 and small-sample-size corrected Akaike information
235 criterion (AIC) were used to assess goodness-of-fit for different regression models.

236 Structural Equation Modeling (SEM) allows testing of multiple separate linear models
237 together into a single causal network, evaluating complex causality between variables by
238 translating the hypothesized causal relationships into a pattern of expected statistical
239 relationships in the data (Jing *et al.* 2015). We used this SEM approach to analyze the relative
240 importance of grazing pressure, soil microbial abundance, plant species richness, and
241 functional diversity, and their interactions on EMF. In our model, we assumed grazing
242 pressure had effects on EMF directly or indirectly by affecting soil microbial abundance,
243 plant species richness, and functional diversity. The standardized coefficient for each path
244 from each model component is shown (Figure 5). Inclusion of these variables in SEM
245 requires us to first test the bivariate relationships between all variables with simple linear
246 regressions to ensure that linear models were appropriate, and then constructed *a priori*
247 model based on the known effects and potential relationships. The chi-square test and its
248 associated p -value were used to adjust the model (good fit when $0 \leq \chi^2 \leq 2$ and $0.05 < p \leq$
249 1.00). The RMSEA statistic (good fit when $0 \leq \text{RMSEA} \leq 0.05$ and $0.10 < p \leq 1.00$) and AIC
250 were used to evaluate the fit of the model (Xu *et al.* 2015). The non-significant pathways
251 were eliminated when significant pathways were left in the final model.

252 **Results**

253 **Relationships between grazing pressure, functional diversity, and plant richness.** Both
254 plant richness (Figure 1a) and functional diversity (Figure 1b) were negatively correlated

255 with grazing pressure. While plant richness showed a weak decline with increasing grazing
256 pressure, FD decreased strongly across the grazing gradient (Fig. 1. a, b). There was no
257 significant relationship between species richness and FD (data not shown, $p = 0.07$, $R^2 =$
258 0.13). While Figure 1b shows the relationship of Mason FD index, we also calculated the
259 community weighted mean of all measured plant functional traits [CWM], which had similar
260 correlations with plant richness and grazing pressure. This loss in FD is due to shifts in the
261 relative abundance of 28 species. Most of dominant species and common species which have
262 major effects on ecosystem processes, decreased their abundance by 30-95% with increasing
263 grazing pressure, with resultant increases in bare ground (Table 1). Of all dominant or
264 common plant species, only *Agropyron cristatum* increased with increased grazing pressure,
265 and *Carex korshinskyi* did not significantly change in relative abundance across the grazing
266 gradient. Of rare plant species, 15 were not present at high grazing pressure, resulting in an
267 overall loss of plant species richness. Linear correlations between edaphic factors and plant
268 richness as well as FD were further tested. Both plant richness and FD were significantly
269 related to plant-available nitrogen but not other factors (except pH, which correlated to plant
270 richness) (Supplementary Table S1).

271 **The effects of grazing on soil microbes.** Grazing significantly reduced the abundance of
272 AM fungi, saprophytic fungi, and soil bacteria (Fig. 2 a-c). Grazing had its weakest effect on
273 AM fungal abundance, which was greatest at moderate grazing pressure (3.0-4.5 sheep ha⁻¹),
274 and only significantly declined at the highest grazing pressure (Fig. 2 a). Increased grazing
275 pressure led to strong linear decreases in saprophytic fungal abundance (Figure 2b), and a
276 curvilinear decrease in bacterial abundance (Fig. 2 c). To further examine relationships

277 between soil microbes and plant richness, aboveground productivity, and FD, we conducted
278 regression analyses with soil microbial abundance as predictors (Supplementary Figs.1, a-i).
279 Plant richness and plant aboveground productivity had weak ($R^2 < 0.17$) positive correlations
280 with AM fungal, saprophytic fungal, and bacterial abundances, in contrast to stronger
281 correlations between FD and AM ($R^2 = 0.31$) and saprophytic fungal abundances ($R^2 = 0.20$)
282 (Supplementary Figs.1, d-h).

283 **The effect of grazing on Ecosystem multifunctionality (EMF).** As hypothesized, high
284 grazing pressure reduced EMF (Fig. 3). EMF was maximized when sheep densities were
285 between 1.5 and 3.0 sheep ha⁻¹, and to maintain 50% of EMF, grazing pressure had to remain
286 below 4.5 sheep ha⁻¹ (Fig. 4). Of the individual functions, plant-available phosphorus did not
287 change in response to the grazing gradient (Supplementary Figure 2d), while plant tissue
288 nitrogen content increased along the grazing gradient (Supplementary figure 2b). All other
289 functions (aboveground net primary production, soil organic carbon, and plant-available
290 nitrogen) decreased with increased grazing (Supplementary Figs. 2, a-e). ANPP sharply
291 declined at the highest grazing pressure, while soil carbon and nitrogen consistently declined
292 with increased grazing pressure.

293 Plant FD had a moderate-strength positive correlation with EMF ($p = 0.01$, $R^2 = 0.20$),
294 while plant richness and soil microbial abundance were weakly positively correlated with
295 EMF (Supplementary Figs. 3 a-c). Structural equation models (SEM) were fitted to infer
296 direct and indirect effects of grazing pressure, soil microbes, plant richness, and FD on EMF
297 (Fig. 5. a, b). Two models were selected based on Chi-square tests ($p > 0.05$), RMSEA ($p >$

298 0.10) and AIC (the least value) statistics. Our SEM indicates grazing pressure directly
299 influenced EMF ($\beta = -0.61$, standardized path coefficients, $p < 0.001$). The indirect effects of
300 soil microbial abundance ($\beta = -0.21$, $p > 0.05$) and plant richness ($\beta = -0.38$, $p > 0.05$) on
301 EMF were not significant (Fig. 5. a). Plant richness, FD, and soil microbial abundance had no
302 interaction or significant direct effect on EMF (Fig. 5. b). When independently assessing the
303 effects of soil fungi or bacteria, on EMF, only the relationship between AM fungal abundance
304 and EMF was significant (Supplementary Figs. 4). However, the direct effect of grazing
305 pressure on EMF was significant ($\beta_a = -0.61$, $p < 0.001$; $\beta_b = -0.48$, $p < 0.01$), and explained
306 22-3% of EMF variation (Fig. 3 and Fig. 5).

307 **Discussion**

308 *Multiple functions are critical to assess ecosystem impacts of grazing pressure.* Moderate
309 grazing pressures (ca. 3.0-4.5 sheep ha⁻¹) have been reported to encourage the greatest plant
310 productivity in semi-arid Eurasian steppe (Liu *et al.* 2015; Li *et al.* 2017). However, setting
311 grazing prescriptions based on only a few ecosystem functions may unintentionally degrade
312 other ecosystem processes (Gordon 1998; MEA 2005; Bennett *et al.* 2009). Our study
313 indicates grazing assessments are more reliable when EMF is tracked, as compared to
314 measuring a single ecosystem function such as ANPP. In our study, moderate grazing
315 pressure (ca. 3.0-4.5 sheep ha⁻¹) did maintain ANPP, but grazing pressure above 3.0 sheep
316 ha⁻¹ directly reduced plant species richness, plant community FD, and most importantly EMF
317 (Figure 1 and supplementary Figure 2). Less intense grazing pressures (1.5-3.0 sheep ha⁻¹)
318 were required to maintain EMF because soil organic carbon and plant-available nitrogen

319 decreased linearly with grazing pressure (Supplementary Figure 2). Similarly, fungal
320 abundance steadily decreased along the grazing gradient (Figure 2a). In contrast, ANPP and
321 bacterial biomass only decreased at high grazing pressures (Figure 2c and supplementary
322 Figure 2a). Therefore, low-intensity grazing is a crucial biotic disturbance that can increase
323 EMF in semi-arid grasslands; however, maintaining 4.5 sheep ha⁻¹ or fewer may be a key
324 grazing pressure tipping point (threshold) for maintaining > 50% EMF in semi-arid
325 grasslands (Figure 3 and Figure 4).

326 ***Plant functional diversity is the strongest indicator of grazing effects on the plant***
327 ***community***. Functional diversity encompasses the range of traits distributed across a plant
328 community, and can be strongly linked to ecosystem properties (Cadotte 2017; Xu *et al.*
329 2018). Plant FD was a more sensitive indicator of grazing effects on the plant community,
330 compared with species richness (Figure 1 and Figure 5b). Plant FD was particularly sensitive
331 to grazing pressure, and likely decreased through both the loss of rare species and decreases
332 in abundance of dominant species (Table 1). Under relatively low grazing pressure, plant
333 communities tend to have a wider variety of complementary traits, resulting in greater FD
334 (Figure 1b). Conversely, functional traits of plant species tend to be more similar under
335 increased grazing pressure, regardless of plant species richness. Diaz *et al.* (2007a) showed
336 grazing can strongly filter plant species by traits, benefiting annual species of short stature,
337 with rosette or stoloniferous architecture. Li *et al.* (2017) demonstrated species with low
338 specific leaf area (SLA) and high leaf nitrogen content (LNC) are negatively affected by
339 intense grazing pressure. Functionally diverse plant communities tend to be resilient to
340 periodic disturbances, thus maintaining ecosystem functions over time (Chapin *et al.* 1997;

341 Diaz & Cabido 2001). Managing plant functional traits in grazed grasslands could regulate
342 species composition for both production and environmental goals, enhancing at least some
343 ecosystem functions and services.

344 ***Links between composition and function.*** Grazing significantly decreases plant FD (and to a
345 lesser extent, richness and biomass), soil organic carbon and microbial biomass, and multiple
346 ecosystem functions (Figure 1, 2 and supplementary figure 2). While there were weak
347 correlations between plant species richness and microbial abundance, microbial abundance
348 and EMF, and between plant species richness and EMF (Supplementary Figure 1 and 3),
349 these were not important drivers of EMF in our Structural Equation Models. EMF was
350 substantially and directly affected by grazing, as opposed to indirectly through the effects of
351 grazing on plant communities or microbial abundance (Figure 5). Similarly, a recent grazing
352 intensity study in drylands showed that decomposition rates were strongly influenced by the
353 direct effects of grazing, not indirectly through grazing effects on FD (Chillo et al. 2017).

354 In summary, grassland management strategies may be flawed when based on monitoring of
355 individual ecosystem functions (Soliveres *et al.* 2016; Stein *et al.* 2016). Our research
356 strongly suggests that the assessment of multiple ecosystem functions is critical to elucidate
357 the optimal grazing thresholds or EMF relationships that ensure the delivery of a suite of
358 ecosystem services critical for sustainable grassland management. Low grazing pressure is
359 required to maintain delivery of multiple functions. Establishing thresholds of grazing to
360 maintain multiple functions is critical for sustainable rangeland management, and can
361 increase prediction accuracy on grassland ecosystem responses to grazing pressure. We need
362 more wide-spread assessment of grazing thresholds for multiple functions across diverse

363 grasslands because many mesic grasslands are predicted to become more arid under a
364 changing climate, and thus are likely to decrease the intensity of grazing that can be
365 sustained.

366

367 **Acknowledgements**

368 We are grateful to all who assisted with collection and processing data over the years. We
369 acknowledge Prof. Friedhelm Taube from Kiel University, Germany, for initial experimental
370 platform construction. This project was supported by National Natural Science Foundation of
371 China (31700389) and (31501996), and Basic research program of Jiangsu province (Natural
372 Science Foundation) -Youth Foundation (BK20160738). We acknowledge the Inner
373 Mongolia Grassland Ecosystem Research Station (IMGERS) of the Chinese Academy of
374 Sciences for providing field facilities and a long-term meteorological dataset. Many thanks
375 are expressed to the anonymous reviewers for their helpful suggestions.

376 **Author contributions**

377 Haiyan Ren, Yongfei Bai, Shuijin Hu, and Yingjun Zhang designed the research; Haiyan Ren,
378 Weiyang Gui, and Gaowen Yang performed the research; Haiyan Ren, Gail W. T. Wilson,
379 Adam B. Cobb, and Valerie Eviner analyzed the data; and all coauthors contributed to the
380 writing of the manuscript.

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583

584 **Table 1.** Relative abundance (RA) of all plant species at low and high grazing pressure in the
 585 Inner Mongolia steppe grassland (lowest grazing level (non zero) /highest grazing level \pm
 586 s.e.m.). Nomenclature follows the editorial committee of Chinese plant records.

Dominant species		Common species	
Latin name	RA (%)	Latin name	RA (%)
<i>Leymus chinensis</i>	(39.25/27.38 \pm 3.58)*	<i>Cleistogenes squarrosa</i>	(10.51/7.32 \pm 1.51)**
<i>Carex korshinskyi</i>	(38.89/36.16 \pm 2.75)	<i>Agropyron cristatum</i>	(3.53/5.00 \pm 1.30)
<i>Stipa grandis</i>	(27.58/12.96 \pm 1.72)**	<i>Achnatherum sibiricum</i>	(2.77/0.14 \pm 0.65)***
Rare species			
Latin name	RA (%)		
<i>Koeleria macrantha</i>	(0.94/0.00 \pm 0.37)***	<i>Potentilla bifurca</i>	(0.08/0.00 \pm 0.07)**
<i>Allium condensatum</i>	(0.04/0.00 \pm 0.02)**	<i>Allium senescens</i>	(0.04/0.00 \pm 0.02)*
<i>Phlomis umbrosa</i>	(0.04/0.00 \pm 0.02)**	<i>Potentilla verticillaris</i>	(0.01/0.00 \pm 0.02)
<i>Adenophora stenanthina</i>	(0.01/0.00 \pm 0.01)	<i>Adenophora gmelinii</i>	(0.01/0.00 \pm 0.01)
<i>Allium tenuissimum</i>	(0.03/0.00 \pm 0.01)*	<i>Poa annua</i>	(0.04/0.00 \pm 0.01)*
<i>Allium anisopodium</i>	(0.02/0.00 \pm 0.01)*	<i>Kochia prostrata</i>	(0.01/0.00 \pm 0.01)
<i>Allium ramosum</i>	(0.01/0.00 \pm 0.01)	<i>Iris tenuifolia</i>	(0.01/0.00 \pm 0.01)
<i>Thalictrum petaloideum</i>	(0.57/0.02 \pm 0.26)**	<i>Potentilla acaulis</i>	(0.05/0.05 \pm 0.02)
<i>Dontostemon micranthus</i>	(0.02/0.02 \pm 0.01)	<i>Axyris amaranthoides</i>	(0.01/0.01 \pm 0.01)
<i>Chenopodium glaucum</i>	(0.01/0.01 \pm 0.01)	<i>Serratula centauroides</i>	(0.01/0.01 \pm 0.01)
<i>Artemisia scoparia</i>	(0.00/0.01 \pm 0.01)	<i>Salsola collina</i>	(0.01/0.04 \pm 0.02)*

587 Note: * means: $0.01 < p < 0.05$, **: $0.001 < p < 0.01$, ***: $p < 0.001$.

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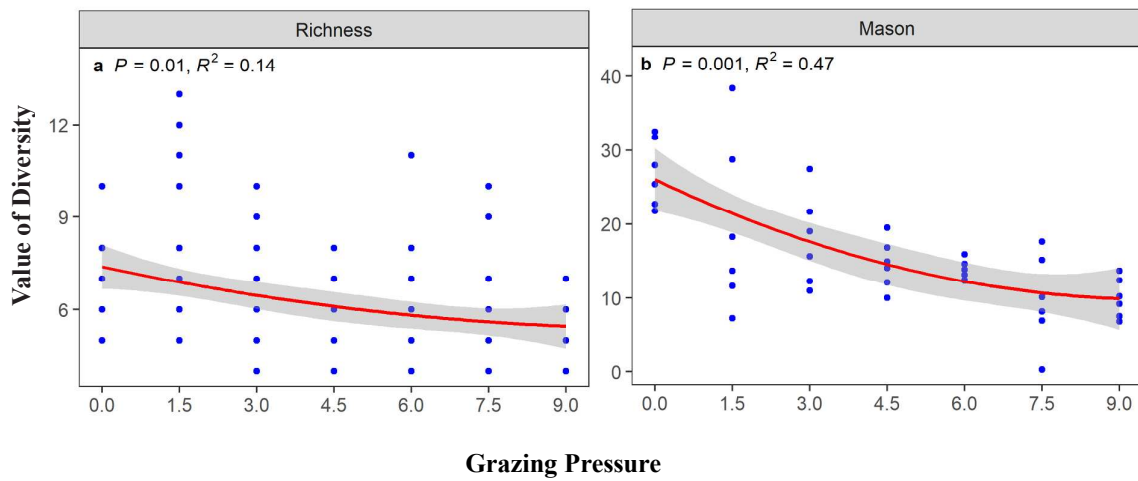
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599 **Figure 1.** Relationship between grazing pressure (number of sheep ha⁻¹) and a) plant species
600 richness or b) plant functional diversity (Mason functional diversity index). Data have been
601 log-transformed. Red lines are fitted lines from OLS regressions. Shaded areas show 95% CI
602 of the fit.

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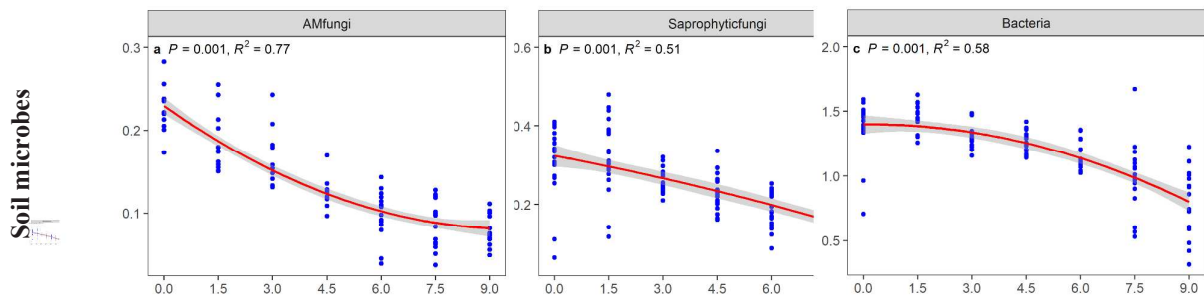
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615 **Grazing Pressure**Figure 2. Relationship between grazing pressure (number of sheep ha^{-1}) and

616 a) arbuscular mycorrhizal (AM) fungal abundance, b) saprophytic fungal abundance, and c)

617 bacterial abundance. Red lines are fitted lines from OLS regressions. Shaded areas show 95%

618 CI of the fit.

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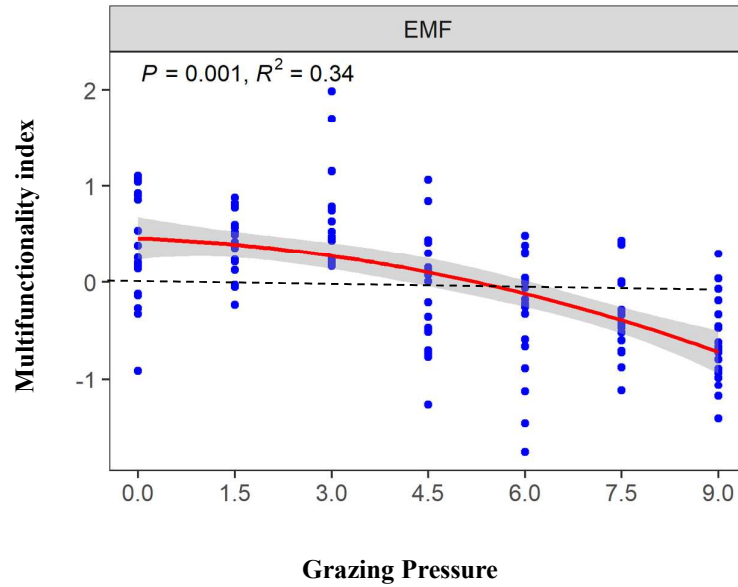
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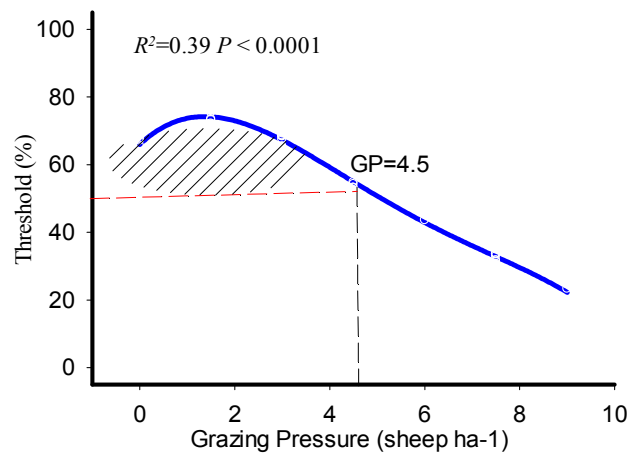


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628 **Figure 3.** Relationship between grazing pressure (number of sheep ha⁻¹) and the
629 multifunctionality index (EMF). Data have been log-transformed. Red lines are the fitted
630 lines from OLS regressions. Shaded areas show the 95% CI of the fit.

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634 **Figure 4.** Relationship between grazing pressure (GP) and ecosystem multifunctionality (EMF)

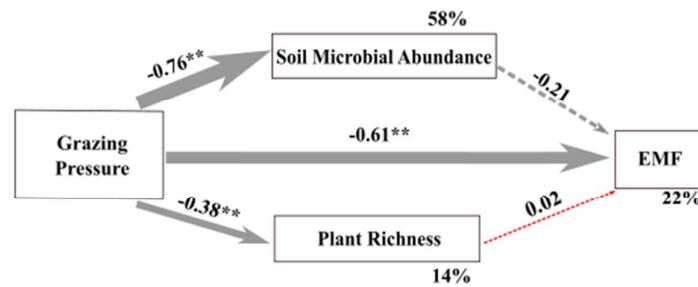
635 with an indication of the 50% EMF threshold level (GP = 4.5, n = 126). The shaded area

636 represents the necessary grazing densities to maintain EMF above 50%.

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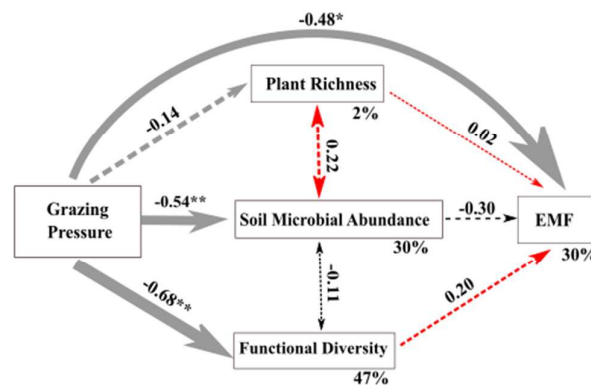
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a. $\chi^2 = 0.45, p = 0.50, AIC = 18.450, RMSEA = 0.00, p = 0.56$

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b. $\chi^2 = 0.03, p = 0.85, AIC = 28.03, RMSEA = 0.00, p = 0.86$

647 **Figure 5.** Structure equation models of grazing pressure, soil microbial abundance, plant
 648 species richness, and functional diversity as predictors of ecosystem multifunctionality
 649 (EMF). Solid red lines represent positive paths ($p < 0.05$, piecewise s.e.m.; ***, $p < 0.001$; **, $p < 0.01$; *, $p < 0.05$), solid gray lines represent negative paths ($p < 0.05$, piecewise s.e.m.)
 650 and dotted gray lines represent non-significant paths ($p > 0.05$, piecewise s.e.m.). Arrow
 651 width is proportional to the strength of the relationship. We report the path coefficients as
 652 standardized effect sizes. Overall fit of piecewise s.e.m. was evaluated using Chi-square test
 653 and RMSEA statistic (if $p > 0.05$, then no paths are missing and the model is a goodfit) and
 654

655 Akaike information criterion (AIC). The proportion of variance explained (R^2) appears
656 alongside response variables in the model.

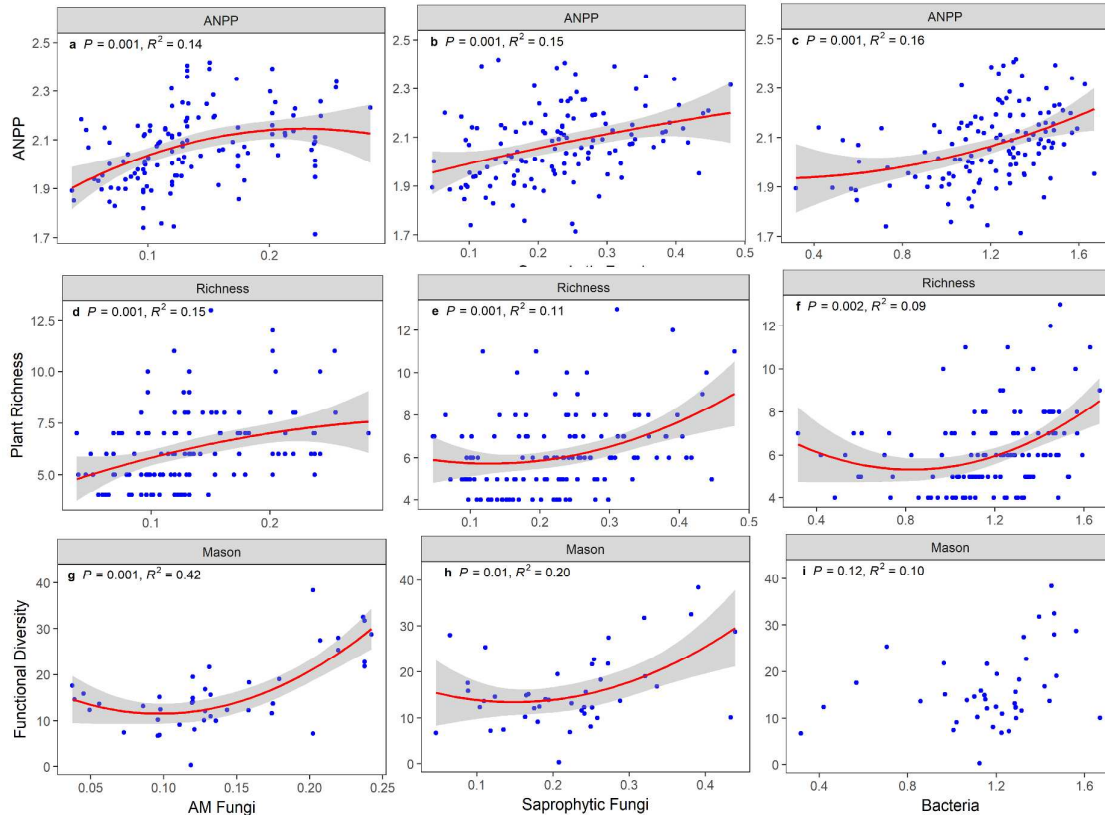
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Table. S1 R²-values statistics from generalized linear models (GLM) between plant richness, functional diversity (FD) and edaphic factors.

Edaphic factors	Plant richness	Edaphic factors	FD
SAN	0.22**	SAN	0.18*
SAP	0.14ns	SAP	0.00ns
SOC	0.07ns	SOC	0.05ns
SM	0.08ns	SM	0.12ns
PH	0.18*	PH	0.03ns

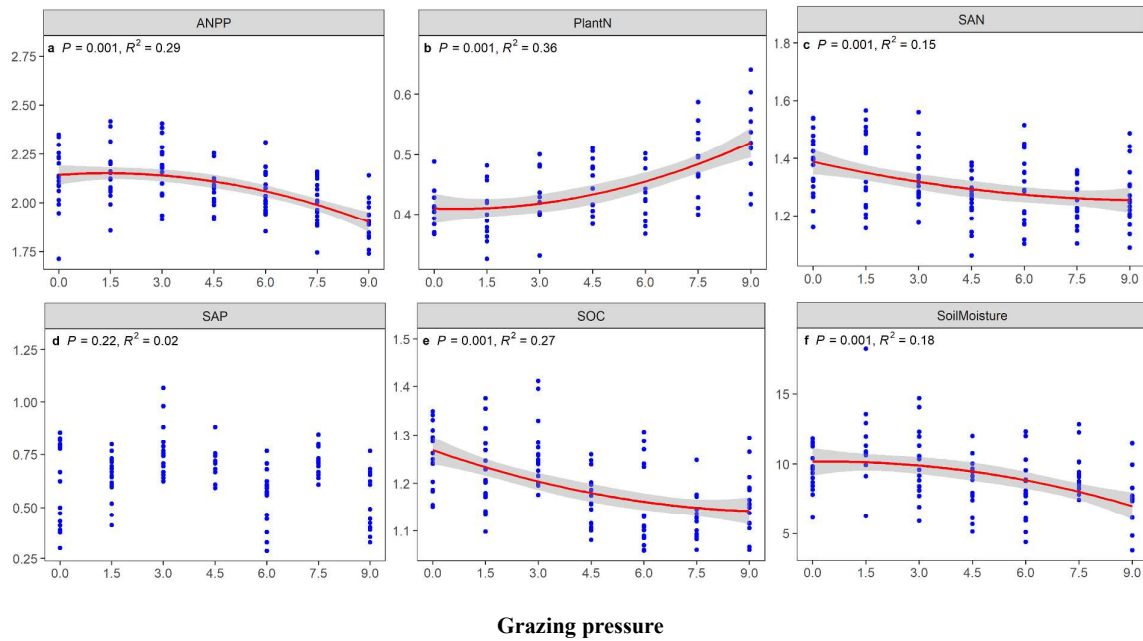
Abbreviations: soil plant-available nitrogen (SAN), soil plant-available phosphorus (SAP), soil organic carbon (SOC), soil moisture (SM).

Stars denote for significance at $p < 0.05$, $p < 0.01$ and $p < 0.001$ probability levels (*, ** and ***, respectively).

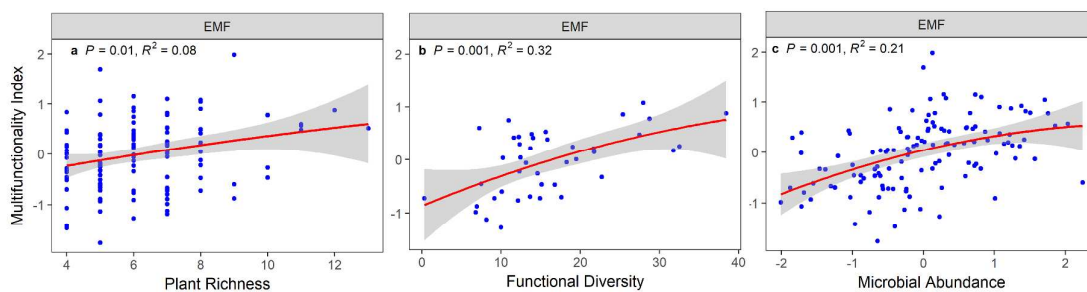


Soil Microbial Abundance

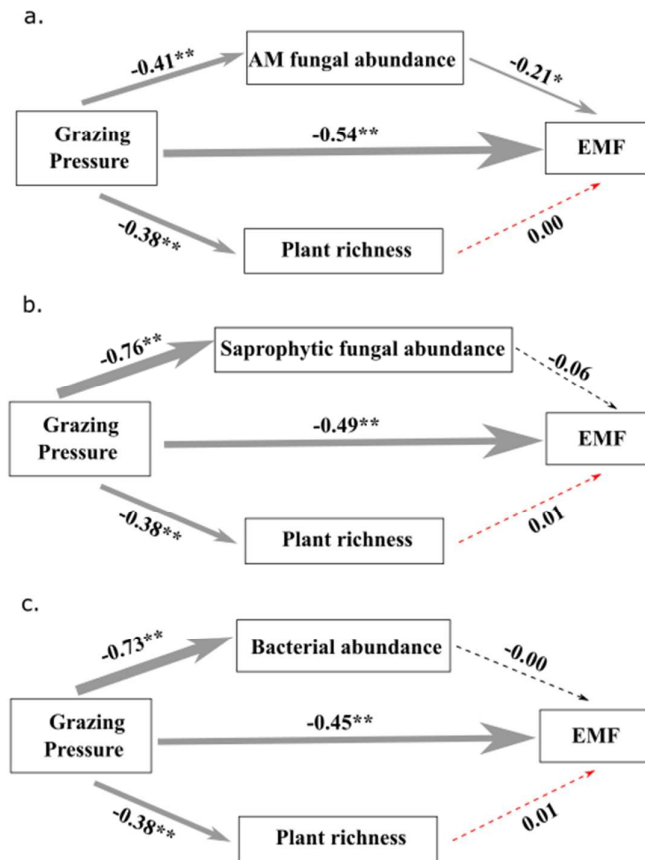
Supplementary Figure 1. Relationships between ANPP (log-transformed) and a) AM fungal abundance, b) saprophytic fungal abundance, and c) bacterial abundance. Relationships between plant species richness and d) AM fungal abundance, e) saprophytic fungal abundance, and f) bacterial abundance. Relationships between functional diversity, and g) AM fungal abundance, h) saprophytic fungal abundance, and i) bacterial abundance. Red lines are the fitted lines from OLS regressions. Shaded areas show the 95% CI of the fit.



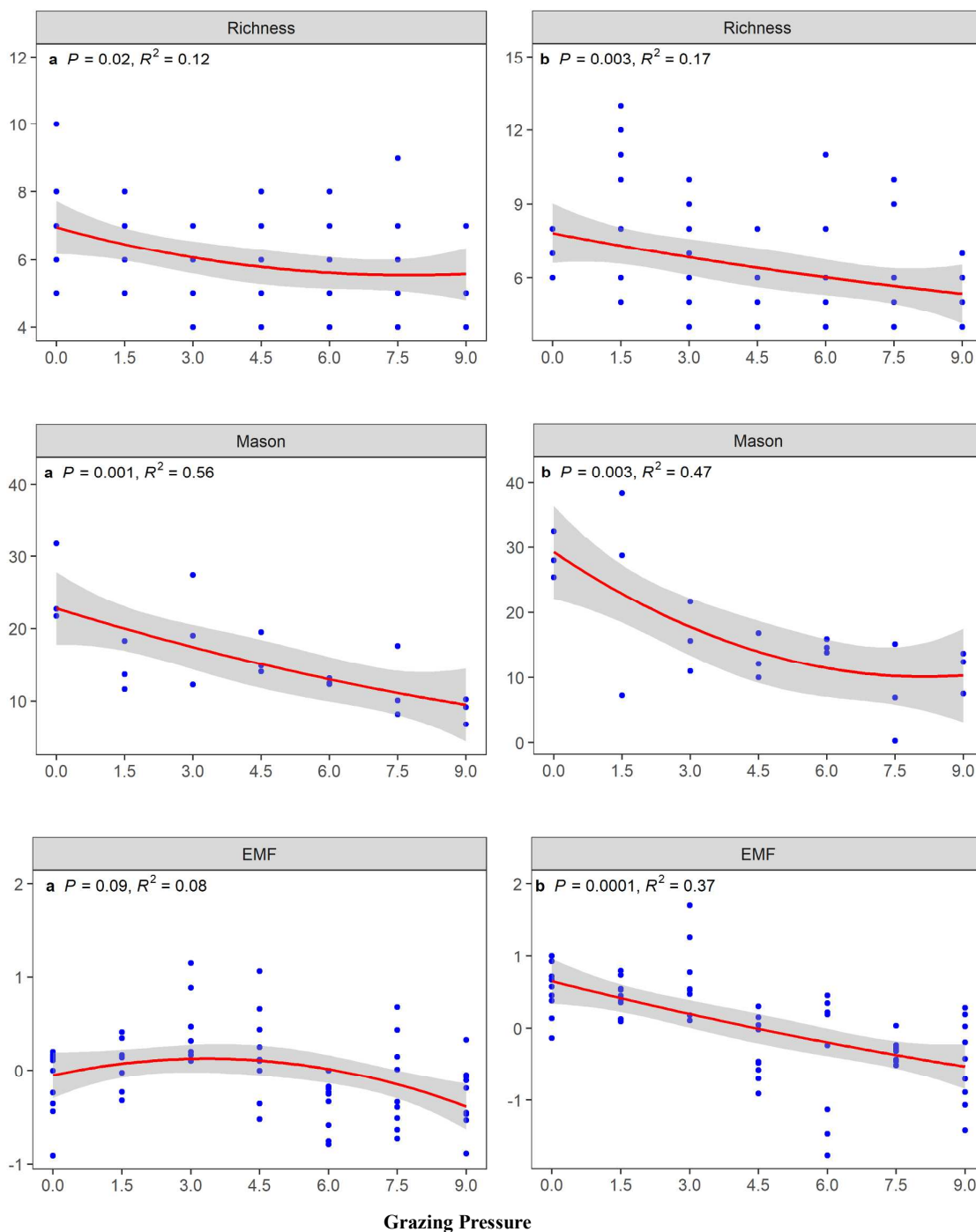
Supplementary Figure 2. Relationship between each component of ecosystem multifunctionality: a) aboveground net primary productivity (ANPP), b) plant-tissue nitrogen content, c) soil plant-available nitrogen, d) soil plant-available phosphorus, e) soil organic carbon, f) soil moisture across grazing densities. Red lines are the fitted lines from OLS regressions. Shaded areas show the 95% CI of the fit.



Supplementary Figure 3. Relationship between ecosystem multifunctionality and a) plant species richness or b) functional diversity (Mason functional diversity index) or c) soil microbial abundance. Red lines are fitted lines from OLS regressions. Shaded areas show 95% CI of the fit.



Supplementary Figure 4. Structure equation models of grazing pressure, soil microbial abundance, and plant species richness as predictors of ecosystem multifunctionality (EMF). Solid red lines represent positive paths ($p < 0.05$, piecewise s.e.m.; ***, $p < 0.001$; **, $p < 0.01$; *, $p < 0.05$), solid gray lines represent negative paths ($p < 0.05$, piecewise s.e.m.) and dotted gray lines represent non-significant paths ($p > 0.05$, piecewise s.e.m.). Arrow width is proportional to the strength of the relationship. We report the path coefficients as standardized effect sizes. a) AM fungal abundance: $\chi^2 = 0.03$, $p = 0.87$, AIC = 18.020, RMSEA = 0.00, $p = 0.89$, b) saprophytic fungal abundance: $\chi^2 = 1.56$, $p = 0.21$, AIC = 19.560, RMSEA = 0.07, $p = 0.28$, c) bacterial abundance: $\chi^2 = 0.23$, $p = 0.63$, AIC = 18.230, RMSEA = 0.00, $p = 0.68$.



Supplementary Figure 5. Relationship between grazing pressure and plant species richness, functional diversity (Mason functional diversity index) and ecosystem multifunctionality (EMF) in (a) sloped system and (b) flat system. Red lines are fitted lines from OLS regressions. Shaded areas show 95% CI of the fit.

GI	System	SAN	SAP	SOC	ANPP	AMF	Bacteria	Fungi
0	1	1.395274	0.498841	1.186269	1.713491	0.23764598	1.33613	0.253892
0	1	1.302205	0.411101	1.286599	2.014605	0.23764598	0.963069	0.272238
0	1	1.27782	0.471517	1.293862	1.991817	0.23764598	1.394619	0.3199
0	1	1.216641	0.388939	1.202915	2.228466	0.22118957	1.488416	0.355654
0	1	1.328253	0.432188	1.249564	2.232878	0.28296075	1.511516	0.40431
0	1	1.26674	0.30103	1.155459	2.254435	0.20482143	1.371767	0.267411
0	1	1.266965	0.388939	1.295623	2.11133	0.23764598	1.39349	0.335231
0	1	1.321619	0.377418	1.249181	2.092791	0.23764598	1.427161	0.340405
0	1	1.15982	0.388939	1.183571	1.94714	0.23764598	1.353139	0.300542
1.5	1	1.204478	0.667692	1.172209	2.058173	0.1585502	1.302944	0.262421
1.5	1	1.418252	0.681335	1.134568	1.991034	0.17419461	1.314365	0.237619
1.5	1	1.289428	0.694563	1.246538	1.856497	0.17484853	1.441217	0.288196
1.5	1	1.237916	0.731988	1.246705	2.060534	0.16180362	1.44322	0.277507
1.5	1	1.318414	0.798362	1.201866	2.119278	0.21289192	1.568569	0.380231
1.5	1	1.230071	0.766442	1.174417	2.198034	0.15501805	1.424963	0.294583
1.5	1	1.156961	0.731988	1.179222	2.072875	0.17937668	1.532394	0.33288
1.5	1	1.191915	0.766442	1.097872	2.036489	0.17419461	1.418438	0.30339
1.5	1	1.298604	0.719867	1.139714	2.151584	0.17419461	1.418438	0.336434
3	1	1.365993	0.766442	1.175775	1.914132	0.15813967	1.287477	0.251201
3	1	1.265285	0.707399	1.241529	2.037692	0.20717053	1.324642	0.272854
3	1	1.281065	0.694563	1.225552	1.933521	0.17928542	1.471846	0.320842
3	1	1.363608	0.694563	1.239594	2.046349	0.18220157	1.282458	0.311924
3	1	1.273635	0.623986	1.244685	2.191339	0.14149476	1.28833	0.210234
3	1	1.31716	0.667692	1.274154	2.285257	0.15386744	1.286806	0.255651
3	1	1.319064	0.667692	1.28393	2.258685	0.2421724	1.271301	0.247016
3	1	1.24032	0.639048	1.198649	2.247949	0.14830983	1.330169	0.242809
3	1	1.175547	0.653606	1.195222	2.192512	0.15399474	1.224461	0.282311
4.5	1	1.258972	0.882076	1.196976	1.927405	0.11997872	1.151395	0.1898
4.5	1	1.128509	0.719867	1.107347	1.914308	0.11997872	1.14674	0.164887
4.5	1	1.245858	0.707399	1.157156	2.003662	0.11997872	1.202883	0.205982
4.5	1	1.220641	0.882076	1.100515	2.020223	0.12532405	1.303377	0.161343
4.5	1	1.277008	0.719867	1.116241	2.086289	0.10867465	1.257287	0.220383
4.5	1	1.329487	0.707399	1.186455	2.239633	0.11997872	1.252986	0.2014
4.5	1	1.373908	0.882076	1.175074	2.117934	0.11997872	1.359996	0.233114
4.5	1	1.297936	0.719867	1.159086	2.124537	0.12572045	1.322412	0.227995
4.5	1	1.141832	0.707399	1.104291	2.110084	0.11997872	1.244274	0.23721
6	1	1.289923	0.766442	1.101784	1.998695	0.09786705	1.199594	0.182112
6	1	1.215738	0.707399	1.110644	2.034682	0.14349666	1.288261	0.240694
6	1	1.352561	0.592196	1.133683	1.94838	0.08604473	1.284074	0.224371
6	1	1.172352	0.557894	1.088975	1.961453	0.08925775	1.105179	0.170019
6	1	1.167772	0.608381	1.10763	1.996033	0.10770302	1.117	0.152314
6	1	1.319614	0.608381	1.084514	1.953115	0.12376525	1.350919	0.23813
6	1	1.34981	0.592196	1.06861	2.030721	0.12989059	1.355513	0.253446
6	1	1.209783	0.575383	1.100922	2.010936	0.07966991	1.039119	0.138343
6	1	1.285368	0.623986	1.130199	2.074158	0.10732485	1.025455	0.14572
7.5	1	1.198465	0.667692	1.089001	1.952179	0.12765602	1.669366	0.433209
7.5	1	1.197225	0.608381	1.084295	1.744919	0.12121249	1.186103	0.24946
7.5	1	1.162696	0.667692	1.12053	1.891537	0.03754745	0.567417	0.088074
7.5	1	1.216973	0.707399	1.2479	2.069125	0.05117245	0.600186	0.099337

7.5	1	1.190673	0.694563	1.146035	1.954371	0.06256284	0.826456	0.100335
7.5	1	1.16214	0.719867	1.176641	2.089505	0.06852505	0.530041	0.094653
7.5	1	1.147082	0.78798	1.128419	2.159176	0.10181609	1.067551	0.189447
7.5	1	1.102797	0.707399	1.129064	2.137291	0.10167543	1.057332	0.151877
7.5	1	1.222472	0.846845	1.14739	2.149804	0.06438979	0.943851	0.146132
9	1	1.135164	0.681335	1.065692	1.893836	0.09622801	0.317613	0.04621
9	1	1.088483	0.766442	1.191012	1.758205	0.11103728	1.020434	0.179872
9	1	1.254949	0.766442	1.212596	1.82007	0.09622801	1.114392	0.163191
9	1	1.255557	0.681335	1.115323	1.740177	0.09622801	0.725824	0.101948
9	1	1.205713	0.608381	1.059243	1.885361	0.09622801	0.58886	0.071584
9	1	1.233372	0.653606	1.156268	1.903741	0.06965243	0.742734	0.09481
9	1	1.271368	0.667692	1.157943	1.891983	0.10287193	1.225881	0.194818
9	1	1.198811	0.608381	1.116156	2.026411	0.10022868	1.11264	0.156433
9	1	1.167719	0.766442	1.186174	1.977358	0.09622801	0.732659	0.125918
0	2	1.425129	0.667692	1.262097	2.085861	0.23668757	1.462494	0.381525
0	2	1.540604	0.78798	1.341391	2.201087	0.21963479	1.462494	0.064404
0	2	1.449455	0.855924	1.331307	2.137523	0.21963479	0.706037	0.11099
0	2	1.460848	0.818409	1.292851	2.33945	0.25522637	1.56799	0.397334
0	2	1.508351	0.798362	1.240231	2.298344	0.20056111	1.452829	0.34207
0	2	1.479009	0.78798	1.310016	2.350778	0.17297391	1.341627	0.356369
0	2	1.538348	0.828096	1.348906	2.059828	0.23469788	1.475922	0.367654
0	2	1.400853	0.777344	1.241686	2.129142	0.2126718	1.485902	0.307898
0	2	1.376871	0.623986	1.152153	2.136594	0.21963479	1.592221	0.410263
1.5	2	1.566343	0.714615	1.314162	2.159938	0.2023475	1.449564	0.390827
1.5	2	1.535127	0.650259	1.314372	2.198171	0.24237718	1.561845	0.439086
1.5	2	1.416883	0.643906	1.228646	2.392064	0.2023475	1.257387	0.118016
1.5	2	1.439779	0.464176	1.281748	2.392903	0.15052539	1.493145	0.31074
1.5	2	1.427643	0.518685	1.142121	2.418008	0.15078725	1.314546	0.142057
1.5	2	1.429374	0.617525	1.268297	2.315446	0.25460758	1.626551	0.479117
1.5	2	1.486356	0.603707	1.376108	2.209475	0.2023475	1.528048	0.446739
1.5	2	1.510279	0.412864	1.35443	2.123982	0.2023475	1.449755	0.385266
1.5	2	1.49536	0.59663	1.206113	2.078602	0.2023475	1.482874	0.417286
3	2	1.331756	0.755258	1.215536	2.177652	0.13130317	1.158754	0.251402
3	2	1.440541	0.743779	1.247914	2.187898	0.13229836	1.287762	0.242838
3	2	1.407447	0.882076	1.329651	2.098505	0.13229836	1.226951	0.241313
3	2	1.560763	1.065618	1.411165	2.359839	0.13229836	1.23704	0.254005
3	2	1.486685	0.98011	1.395169	2.385904	0.13229836	1.202582	0.227103
3	2	1.339047	0.808501	1.257293	2.406245	0.13229836	1.30394	0.238687
3	2	1.403091	0.755258	1.215224	2.249345	0.13329127	1.2397	0.233253
3	2	1.284813	0.743779	1.259734	2.155973	0.13229836	1.484037	0.297949
3	2	1.303805	0.78798	1.201136	2.19064	0.13229836	1.346729	0.278565
4.5	2	1.28649	0.681335	1.102729	2.074938	0.12849221	1.418438	0.336434
4.5	2	1.190173	0.667692	1.114857	2.007463	0.12849221	1.158221	0.175034
4.5	2	1.062363	0.592196	1.080549	1.988381	0.13577522	1.289678	0.25826
4.5	2	1.269638	0.681335	1.201641	2.091127	0.16976849	1.372697	0.293069
4.5	2	1.230236	0.608381	1.146184	2.11789	0.0966392	1.169173	0.209871
4.5	2	1.221115	0.743779	1.2389	2.254064	0.12849221	1.233874	0.264291
4.5	2	1.357863	0.755258	1.259741	2.125026	0.11683134	1.143211	0.206952
4.5	2	1.384471	0.719867	1.246832	2.053194	0.11998219	1.194522	0.224096
4.5	2	1.226027	0.592196	1.105928	2.088632	0.12849221	1.295378	0.279659

6	2	1.437937	0.603707	1.240229	2.311415	0.11943766	1.068302	0.194441
6	2	1.45162	0.464176	1.30671	1.85187	0.03902912	1.101189	0.123737
6	2	1.515322	0.574679	1.23395	2.184578	0.04512959	1.13047	0.088263
6	2	1.4485	0.378888	1.287055	2.157583	0.09172084	1.10931	0.179978
6	2	1.380844	0.680698	1.269885	2.143697	0.11434869	1.206449	0.234793
6	2	1.444956	0.567109	1.288358	2.147367	0.11434869	1.094542	0.234793
6	2	1.116189	0.287443	1.057629	1.941362	0.09172084	1.031045	0.165009
6	2	1.185211	0.444373	1.059401	2.092931	0.11005241	1.086637	0.229899
6	2	1.102156	0.329011	1.130337	1.979321	0.09172084	1.089436	0.219851
7.5	2	1.227192	0.719867	1.100395	2.123263	0.09724567	0.968732	0.167459
7.5	2	1.254801	0.653606	1.05935	1.963364	0.09724567	1.224232	0.222199
7.5	2	1.314736	0.719867	1.081849	1.880013	0.11862459	1.124236	0.20768
7.5	2	1.353744	0.707399	1.12788	2.010944	0.09724567	1.060635	0.194936
7.5	2	1.358007	0.731988	1.137144	1.908764	0.09912707	1.010832	0.164624
7.5	2	1.227122	0.694563	1.093234	2.002252	0.09724567	1.098175	0.176415
7.5	2	1.292733	0.798362	1.172839	1.952356	0.12250594	1.010832	0.114961
7.5	2	1.344434	0.731988	1.11994	1.931407	0.05919629	0.983504	0.138426
7.5	2	1.29028	0.639048	1.126894	2.040325	0.0836568	0.901629	0.169636
9	2	1.488034	0.492285	1.294272	2.141157	0.04935378	0.420388	0.103876
9	2	1.442579	0.444373	1.265004	1.829132	0.0728373	1.007349	0.134735
9	2	1.425005	0.401831	1.150244	1.938853	0.05587306	0.85827	0.109221
9	2	1.351613	0.624272	1.165407	1.845931	0.06852505	0.594003	0.092738
9	2	1.206693	0.354665	1.187976	1.900003	0.07556636	0.978065	0.180352
9	2	1.247024	0.444373	1.189471	2.00107	0.06852505	0.60349	0.048375
9	2	1.308594	0.423623	1.156443	1.936815	0.08155517	0.922135	0.105761
9	2	1.303689	0.390511	1.138141	1.89603	0.06181701	0.48282	0.097095
9	2	1.220696	0.329011	1.1058	1.898999	0.08154813	0.912853	0.130143

Microbial index	Plant N	Richness	Moisture	PH	EMF	Mason
0.810306967	0.428527	5	8.456513	7.05	-0.324887	22.71
0.435986775	0.488438	6	9.691756	7.12	0.1480468	21.76
1.137145292	0.413818	7	11.56808	7.4	0.1723901	31.77
1.295871979	0.412015	8	8.700251	7.39	-0.121094	NA
1.860193608	0.366943	7	8.700251	8.16	0.532705	NA
0.7209549	0.38347	10	9.623736	7.27	-0.268215	NA
1.195477272	NA	6	9.823949	7.09	0.201229	NA
1.255849414	NA	6	8.184455	8.22	0.1417794	NA
1.012228712	NA	6	7.828354	8.07	-0.90914	NA
0.359131257	0.400223	7	10.73698	8	-0.04964	18.28
0.364191505	0.482134	8	6.275834	7.18	0.4189267	11.63
0.716315629	0.463159	7	18.27747	7.06	0.2118911	13.69
0.603754241	0.457319	8	10.73698	6.95	0.4912155	NA
1.440656395	0.419679	7	6.275834	8.01	0.8213713	NA
0.610260023	0.42307	8	18.27747	7.15	0.4984331	NA
1.024661549	NA	5	10.73698	6.89	-0.0268	NA
0.744571231	NA	6	6.275834	7.05	-0.234038	NA
0.873205658	NA	6	18.27747	7.1	0.4116075	NA
0.294666259	0.429323	7	12.00332	8.23	0.4341047	12.25
0.699046444	0.480157	6	10.55013	7.54	0.4714804	27.45
0.90494545	0.50091	7	14.69306	7.04	0.2425025	19.04
0.660343647	0.482838	7	14.06681	8.04	0.7838649	NA
0.042623568	0.420391	6	7.73927	7.39	0.6317314	NA
0.287164323	0.420823	6	12.33942	6.94	1.1552362	NA
0.731533863	NA	6	10.5634	7	1.1515206	NA
0.257738709	NA	4	10.54189	6.91	0.4706136	NA
0.317174565	NA	5	11.28735	6.8	0.1660757	NA
-0.321509687	0.415914	6	9.588334	7.9	0.412896	14.06
-0.424055618	0.493564	8	7.43788	6.81	-0.726502	14.90
-0.196999345	0.479491	7	9.660303	7.08	0.0063513	19.49
-0.220643536	0.511291	4	12.03414	7.9	0.0707825	NA
-0.139496755	0.396872	6	9.159555	6.81	0.1246931	NA
-0.154973322	0.405751	4	9.140186	7.08	0.8400955	NA
0.096340534	NA	5	9.525423	7.9	1.0629157	NA
0.063789674	NA	4	10.76376	6.81	0.4367304	NA
-0.025983303	NA	4	6.129454	7.08	-0.356528	NA
-0.41817838	0.388105	7	7.243843	7.03	0.0400499	12.44
0.172369482	0.437139	8	4.368322	6.79	-0.223008	12.32
-0.219203881	0.492962	6	5.936993	7.21	-0.05944	13.15
-0.626468499	0.501862	7	10.29712	7.12	-0.885318	NA
-0.577555033	0.442837	5	8.908016	7	-0.662575	NA
0.126312199	0.476885	5	7.83271	8.2	-0.325579	NA
0.225862571	NA	5	6.109902	6.96	-0.176329	NA
-0.882612374	NA	5	8.794801	6.9	-0.586146	NA
-0.714725974	NA	4	7.930716	6.82	-0.003113	NA
2.243919227	0.428817	9	9.42545	6.88	-0.598288	10.10
-0.040862236	0.58732	7	7.829738	6.81	-1.121625	8.16
-1.878733716	0.526453	7	8.241745	6.86	-0.704335	17.60
-1.71912544	0.536542	5	8.613644	6.86	0.3932109	NA

-1.380850295	0.467664	6	12.8704	6.94	-0.339469	NA	
-1.72360311	0.466089	6	8.279931	7.03	-0.012905	NA	
-0.525185558	NA	4	9.146504	6.93	0.0072236	NA	
-0.684438642	NA	4	12.29601	6.87	-0.338012	NA	
-1.052032399	NA	4	10.15277	7.13	0.4343868	NA	
-2.010228781	0.434139	7	4.838945	6.94	-0.984504		6.79
-0.566908741	0.641682	7	6.161221	6.81	-0.723224		9.17
-0.602848383	0.555358	5	3.753392	6.84	0.0351729		10.24
-1.305517989	0.603024	7	4.838945	7.46	-0.667083	NA	
-1.587364246	0.519696	5	6.161221	7.2	-0.933491	NA	
-1.462524412	0.510916	5	3.753392	6.96	-0.329975	NA	
-0.309167446	NA	5	4.838945	7.05	-0.189941	NA	
-0.608753911	NA	4	6.161221	6.94	-0.468326	NA	
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0.318970195	0.41096	8	11.23835	8.13	1.0760396		27.96
-0.473831718	0.43968	6	9.363605	8.13	0.8599408		25.38
1.710302396	0.404981	8	9.013975	8.18	1.0448977	NA	
1.073062066	0.369889	8	10.399	8.33	0.9036025	NA	
0.818844205	0.404688	7	11.86802	8.28	1.10162	NA	
1.382116578	NA	6	9.360016	8.28	0.9265855	NA	
1.076937999	NA	6	11.44597	8.23	0.380666	NA	
1.573276821	NA	6	6.17305	7.25	-0.135311	NA	
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1.759534164	0.389194	10	13.58604	8.22	0.7768131		28.75
0.132502081	0.32645	11	10.80154	8.25	0.5968011		7.26
0.731730865	0.371546	13	11.28123	8.34	0.5181101	NA	
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1.507784841	NA	5	9.919435	8.21	0.7915002	NA	
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1.358560619	NA	6	10.6187	8.33	0.1247649	NA	
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0.155380355	0.429113	7	9.174018	8.27	0.5235187		15.63
0.074990049	0.419679	6	8.83533	8.19	0.7451154		10.97
0.12764315	0.400125	9	10.34366	8.17	1.9849951	NA	
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0.071162916	NA	6	8.410845	7.87	0.4719047	NA	
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0.341573987	NA	5	8.128419	7.64	0.1821737	NA	
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