

Consistent ecosystem functional response across precipitation extremes in a sagebrush steppe

Andrew T. Tredennick¹, Andrew R. Kleinhesselink², Bret Taylor³, and Peter B. Adler¹

¹Department of Wildland Resources and the Ecology Center, Utah State University, Logan, Utah 84322

²Department of Ecology and Evolutionary Biology, University of California, Los Angeles, Los Angeles, California 90095

³United States Department of Agriculture, Agriculture Research Station, U.S. Sheep Experiment Station, Dubois, Idaho 83423

Corresponding author:
Andrew T. Tredennick¹

Email address: atredenn@gmail.com

ABSTRACT

Background. Precipitation is predicted to become more variable in the western United States, meaning years of above and below average precipitation will become more common. Periods of extreme precipitation are major drivers of ecosystem functioning in water limited grasslands, but how ecosystems respond to precipitation may change as the duration of above and below average periods increases. Changes in ecosystem functional response could reflect compensatory changes in species composition or species reaching physiological thresholds at extreme precipitation levels.

Methods. We conducted a five year precipitation manipulation experiment in a sagebrush steppe ecosystem in Idaho, US. We used drought and irrigation treatments (approximately 50% decrease/increase) to investigate whether ecosystem functional response remains consistent at extreme precipitation levels. We recorded data on aboveground net primary productivity (ANPP), species abundance, and soil moisture. We fit a random intercept, random slope model to determine if the relationship between ANPP and soil moisture differed among treatments. We used nondimensional multivariate scaling to quantify community composition over the five years.

Results. Ecosystem functional response, defined as the relationship between soil moisture and ANPP, did not vary across drought, control, and irrigation treatments. ANPP responded positively to available soil moisture, but the response was weak and uncertain. There was also no evidence that treatment effects grew over time. Plant community composition was remarkably stable over the course of the experiment and did not differ among treatments.

Discussion. At least in the short-term, ecosystem functional response and community composition in this sagebrush steppe system is consistent, even at extreme precipitation levels. The similarity of ecosystem functional response across treatments was not due to compensatory shifts at the plant community level, but instead reflects the consistent responses of individual species. Such consistency may be due to bet-hedging strategies where species respond weakly to soil moisture to ensure low variance of long-term success.

1 INTRODUCTION

At a given site, the functional response of aboveground net primary productivity (ANPP) to water availability (e.g., soil moisture) can be characterized by fitting a model to historical observations of ANPP and soil moisture. However, the fitted functional response may provide an incomplete picture because future conditions are likely to be outside the historical range of variability (Smith, 2011). For example, historical trends may underestimate the potential for the soil moisture-ANPP relationship to saturate if

46 soil moisture is pushed far beyond typical levels. Saturating relationships are actually common (Hsu et al.,
47 2012; Gherardi and Sala, 2015b), perhaps because other resources, like nitrogen, become more limiting
48 in wet years than dry years. Knowing the curvature of the soil moisture-ANPP relationship at extreme
49 precipitation levels is critical for understanding how ecosystems will respond to chronic alterations in
50 water availability.

51 Another problem with relying on historical ecosystem functional responses is that they are not static.
52 Changes in species identities and abundances can alter an ecosystem's functional response to water
53 availability because different species have different physiological thresholds for producing biomass.
54 Smith et al. (2009) introduced the 'Hierarchical Response Framework' (HRF) for understanding the
55 interplay of community composition and ecosystem functioning in response to resource manipulations
56 over time. In the near term, ecosystem functioning such as ANPP will reflect the physiological responses
57 of individual species to the manipulated resource level. For example, ANPP may decline under simulated
58 drought because the initial community consisted of drought-intolerant species (Hoover et al., 2014). Over
59 longer time spans, ecosystem functioning may recover as new species colonize or initial species reorder
60 in relative abundance. For example, ANPP may initially decline, but eventually rise back to pre-treatment
61 levels once drought-tolerant species become more abundant and compensate for drought-intolerant species
62 (Hoover et al., 2014). It is also possible that ecosystem functioning shifts to a new mean state, reflecting
63 the suite of species in the new community (Knapp et al., 2012).

64 Manipulating potentially limiting resources, like precipitation, offers a route to understanding how
65 ecosystems will respond to resource levels that fall outside the historical range of variability (Avolio et al.,
66 2015; Gherardi and Sala, 2015a; Knapp et al., 2017). Altering the amount of precipitation over many years
67 should provide insight into the time scales at which water-limited ecosystems respond to chronic resource
68 alteration. Following the HRF, we propose four alternative predictions for the effect of precipitation
69 manipulation on the ecosystem functional response to soil moisture, that is, the soil moisture-ANPP
70 relationship (Fig. 1). The four predictions are based on possible outcomes at the community (e.g.,
71 community composition) and ecosystem (e.g., soil moisture-ANPP regression) levels.

72 First, altered precipitation changes neither ecosystem functional response nor community composition
73 (Fig. 1, top left). In this case, changes in ANPP simply follow the soil moisture-ANPP relationship under
74 ambient conditions. This corresponds to the early phases of the HRF, where ecosystem response is due to
75 the physiological responses of individual species. Second, the ecosystem functional response changes but
76 community composition remains the same (Fig. 1, top right). A saturating soil moisture-ANPP response
77 fits this scenario, where individual species hit physiological thresholds or are limited by some other
78 resource. Third, the ecosystem functional response is consistent but underlying community composition
79 changes (Fig. 1, bottom left). In this case, changes in species' identities or abundances occur in response to
80 altered precipitation levels and species more suited to the new conditions compensate for reduced function
81 of initial species. Fourth, and last, both ecosystem functional response and community composition
82 change (Fig. 1, bottom right). New species, or newly abundant species, with different physiological
83 responses completely reshape the ecosystem functional response.

84 All four outcomes are possible in any given ecosystem, but the time scales at which the different
85 scenarios play out likely differ (Smith et al., 2009; Wilcox et al., 2016; Knapp et al., 2017). Thus, our
86 task is not to test the validity of the HRF, but rather to amass information on how quickly ecosystem
87 functional responses change in different ecosystems. Likewise, we need to understand whether changes at
88 the ecosystem level are driven by community level changes or individual level responses.

89 To that end, here we report the results of a five-year precipitation manipulation experiment in a
90 sagebrush steppe grassland. We imposed drought and irrigation treatments (approximately $\pm 50\%$) and
91 measured ecosystem (ANPP) and community (species composition) responses. We focus on how the
92 drought and irrigation treatments affect the relationship between available soil moisture and ANPP, and if
93 community dynamics underlie the ecosystem responses. In particular, we are interested in the consistency
94 of the soil moisture-ANPP relationship among treatments. Is the relationship steeper under the drought
95 treatment, at low soil moisture? Does the relationship saturate under the irrigation treatment, at high soil
96 moisture? To answer these questions we fit a random intercept, random slope model to test whether the
97 regressions differed among treatments. We also analyzed community composition over time, allowing us
98 to place our experimental results within the framework of the HRF and our competing predictions (Fig.
99 1).

100 2 METHODS

101 2.1 Study Area

102 We conducted our precipitation manipulation experiment in a sagebrush steppe community at the United
103 States Sheep Experimental Station (USSES) near Dubois, Idaho (44.2° N, 112.1° W), 1500 m above
104 sea level. The plant community is dominated by the shrub *Artemisia tripartita* and three perennial
105 bunchgrasses, *Pseudoroegneria spicata*, *Poa secunda*, and *Hesperostipa comata*. During the period of our
106 experiment (2011 – 2015), average mean annual precipitation was 265 mm year⁻¹ and mean monthly
107 temperature ranged from -5.2°C in January to 21.8°C in July. Between 1926 and 1932, range scientists
108 at the USSES established 26 permanent 1 m² quadrats to track vegetation change over time. In 2007,
109 we (well, one of us [P. Adler]) relocated 14 of the original quadrats, six of which were inside a large,
110 permanent livestock enclosure. We use these six plots as control plots that have received no treatment,
111 just ambient precipitation, in the experiment described below.

112 2.2 Precipitation Experiment

113 In spring 2011, we (well, two of us [A. Kleinhesselink and P. Adler]) established 16 new 1 m² plots located
114 in the same enclosure as the six control plots. We avoided areas on steep hill slopes, areas with greater
115 than 20% cover of bare rock, and areas with greater than 10% cover of the shrubs *Purshia tridentata*
116 and/or *Amelanchier utahensis*. We established the new plots in pairs and randomly assigned each plot in a
117 pair to receive a “drought” or “irrigation” treatment.

118 Drought and irrigation treatments were designed to decrease and increase the amount of ambient
119 precipitation by 50%, respectively. To achieve this, we used a system of rain-out shelters and automatic
120 irrigation (Gherardi and Sala, 2013). The rain-out shelters consisted of transparent acrylic shingles 1-1.5
121 m above the ground that covered an area of 2.5 × 2 m. The shingles intercepted approximately 50% of
122 incoming rainfall, which was channeled into 75 liter containers. Captured rainfall was then pumped out
123 of the containers and sprayed on to the adjacent irrigation plot via two suspended sprinklers. Pumping
124 was triggered by float switches once water levels reached about 20 liters. We disconnected the irrigation
125 pumps each fall and reconnected them, often with difficulty, each spring. The rain-out shelters remained
126 in place throughout the year.

127 We monitored soil moisture in four of the drought-irrigation pairs using Decagon Devices (Pullman,
128 Washington) 5TM and EC-5 soil moisture sensors. We installed four sensors around the edges of each plot,
129 two at 5 cm soil depth and two at 25 cm soil depth. We also installed four sensors in areas nearby the four
130 selected plot pairs to measure ambient soil moisture at the same depths. Soil moisture measurements were
131 automatically logged every four hours. We coupled this temporally intensive soil moisture sampling with
132 spatially extensive readings taken at six points within all 16 plots and associated ambient measurement
133 areas. These snapshot data were collected on 06/06/2012, 04/29/2015, 05/07/2015, 06/09/2015, and
134 05/10/2016 using a handheld EC-5 sensor.

135 Analyzing the response to experimental treatments was complicated by the fact that we did not directly
136 monitor soil moisture in each plot on each day of the experiment. Only a subset of plots were equipped
137 with soil moisture sensors, and within those plots, one or more of the sensors frequently failed to collect
138 data. To remedy these problems, and to produce average daily soil moisture values for the ambient,
139 drought, and irrigation conditions, we used a statistical model to predict the average treatment effects on
140 soil moisture during the course of the experiment.

141 We first averaged the observed soil moisture for each day and within each plot. Then we standardized
142 the averages within each plot group by subtracting the average ambient soil moisture in that plot group
143 and dividing by the standard deviation of the ambient soil moisture in that plot group. We then found the
144 difference between the standardized ambient soil moisture and the standardized drought and irrigation
145 soil moisture within each plot group. These transformations ensured that the treatment effects in each plot
146 were appropriately scaled by the local ambient conditions within each plot group.

147 We then modeled the daily deviation from ambient conditions of the drought and irrigation treatments
148 using a linear mixed effects model with independent variables for treatment, season (winter, spring,
149 summer, fall), rainfall, and all two-way interactions. Rainy days were defined as any day in which
150 precipitation was recorded and average temperature was above 3°C. The day immediately following
151 rainfall was also classified as rainy. We fit the model using the ‘lme4::lmer()’ function (Bates et al., 2015)

in R (R Core Team, 2016), with random effects for plot group and date. We weighted observations by the number of unique sensors or spot measurements that were taken in each plot on that day. We then used the model to predict the average daily soil moisture in the treated plots based on the average daily ambient soil moisture. We could only predict soil moisture in the treated plots on days for which we took at least one ambient soil moisture measurement.

2.3 Data Collection

We estimated aboveground net primary productivity (ANPP) using a radiometer to relate ground reflectance to plant biomass (see Byrne et al., 2011, for a review). We recorded ground reflectance at four wavelengths, two associated with red reflectance (626 nm and 652 nm) and two associated with near-infrared reflectance (875 nm and 859 nm). At each plot in each year, we took four readings of ground reflectances at the above wavelengths. We also took readings in 12 (2015), 15 (2012, 2013, 2014), or 16 (2016) calibration plots adjacent to the experimental site, in which we harvested all aboveground biomass, dried it to a constant weight at 60°C, and weighed it to estimate ANPP.

For each plot and year, we averaged the four readings for each wavelength and then calculated a greenness index based on the same bands used to calculate NDVI using the MODIS and AVHRR bands for NDVI. To convert the greenness index to ANPP we regressed the greenness index against the dry biomass weight from the ten calibration plots. We fit regressions to MODIS-based index and AVHRR-based index for each year and retained the regression with the better fit. Using the best regression equation for each year, we predicted ANPP (Appendix 1).

Species composition data came from two sources: yearly census maps for each plot made using a pantograph (Hill, 1920) and yearly counts of annual species in each plot. From those maps, we have data on the density of all annuals and perennials forbs, basal cover of perennial grasses, and canopy cover of shrubs. We made a large plot-treatment-year by species matrix, where columns were filled with either basal cover or density, depending on the measurement made for the particular species. So we could analyze the different types of data together, we standardized the values in each column. This puts all abundance values on the same scale, meaning that common and rare species are weighted equally. Nonetheless, if we assume that rare species will respond more than common ones, then this approach is anti-conservative for detecting community change. This means that our approach is biased toward detecting compositional changes.

2.4 Data Analysis

Our main goal was to test whether the relationship between ANPP and soil moisture differed among the drought, control, and irrigation treatments. Based on our own observations and previous work at our study site (Blaisdell, 1958; Dalgleish et al., 2011; Adler et al., 2012), we chose to use cumulative volumetric water content from March through June as our metric of soil moisture (hereafter referred to simply as ‘soil moisture’). To achieve this goal, we fit a multi-level random intercept and random slope regression with $\log(\text{ANPP})$ as the response variable and soil moisture as the sole predictor. We log-transformed ANPP to account for heteroscedasticity. We fit the model under a Bayesian framework, allowing us to test for treatment differences by comparing the posterior distributions of the treatment-level coefficients (e.g., Tredennick et al., 2013). Both $\log(\text{ANPP})$ and soil moisture were standardized to have mean 0 and unit variance before fitting the model [i.e., $(x_i - \bar{x})/\sigma_x$].

Our model is defined as follows:

$$\mu_{j(k(t))} = \beta_{0,j(k)} + \beta_{1,j(k)} x_{k(t)} + \gamma_t, \quad (1)$$

$$y_{j(k(t))} \sim \text{Normal}(\mu_{j(k(t))}, \sigma_k^2), \quad (2)$$

where $\mu_{j(k(t))}$ is the deterministic prediction from the regression model for plot j associated with treatment k in year t , $\beta_{0,j(k)}$ is the intercept for plot j associated with treatment k , $\beta_{1,j(k)}$ is the slope term for the effect of soil moisture for plot j associated with treatment k , γ_t is the intercept offset for year t , and σ_k^2 is the process variance for treatment k . Data include the standardized $\log(\text{ANPP})$ observations ($y_{j(k(t))}$) and soil moisture ($x_{k(t)}$).

The intercept and slope terms are modeled hierarchically to account for the non-independence of observations across years within plots and to allow us to test the hypothesis that our treatments alter the

ANPP-soil moisture relationship. Thus, our multi-level model has three grouping levels for coefficients, representing the nested structure of the data: (i) overall coefficients, (ii) treatment coefficients, and (iii) plot coefficients. Each subsequent level is drawn from the distribution of coefficients at the previous level, meaning plot-level coefficients are drawn from treatment-level coefficients, which are drawn from overall coefficients. We also include a covariance structure among the intercept and slope at each level. Formally, our hierarchical structure is as follows, where we drop the intercept (0) and slope (1) subscripts and instead refer to a vector of coefficients, β :

$$\beta_{j(k)} \sim \text{MVN}(\beta_k, \Sigma(k)), \quad (3)$$

$$\beta_k \sim \text{MVN}(\beta, \Sigma), \quad (4)$$

$$\beta \sim \text{Normal}(0, 1), \quad (5)$$

where $\beta_{j(k)}$ is the vector of regression coefficients (intercept and slope) for plot j associated with treatment k , β_k is the vector of coefficients for each treatment, and β is the vector of overall coefficients. The plot- and treatment-level coefficients are drawn from multivariate normal distributions with covariance matrix Σ . For the plot-level coefficients, each treatment has its own variance-covariance matrix (i.e., $\Sigma(k)$). The overall coefficients are drawn from a normal prior with mean 0 and standard deviation 1. The random year effects (γ) are drawn from a normal prior with mean 0 and standard deviation σ_{year} , which was drawn from a weibull distribution. A full description of model is in Appendix 2.

We fit the model using a Bayesian approach, obtaining posterior estimates of all unknowns via the No-U-Turn Hamiltonian Monte Carlo sampler in Stan (Stan Development Team, 2016b). We used the R package ‘rstan’ (Stan Development Team, 2016a) to link R (R Core Team, 2016) to Stan. We obtained samples from the posterior distribution for all model parameters from four parallel MCMC chains run for 10,000 iterations, saving every 10th sample. Trace plots of all parameters were visually inspected to ensure well-mixed chains and convergence. We also made sure all scale reduction factors (\hat{R}) were less than 1.1.

To see if community composition differed among treatments through time, we used non-dimensional multivariate scaling (NMDS) based on Bray-Curtis distances. For each year of the experiment, we first calculated Bray-Curtis distances among all plots, and then extracted those distances for use in the NMDS. Because we standardized species’ abundances, some values were negative, which is not allowed for calculating Bray-Curtis distances. We simply added ‘2’ to each abundance value to ensure all values were greater than zero. We plotted the first two axes of NMDS scores to see if community composition overlapped, or not, among treatments in each year. We used the ‘metaMDS()’ function in the R package ‘vegan’ (Oksanen, 2016) to calculate Bray-Curtis distances and then to run the NMDS analysis. We used the ‘vegan::adonis()’ function (Oksanen, 2016) to perform permutational multivariate analysis of variance to test whether treatment plots formed distinct groupings. To test whether treatment plots were equally dispersed, or not, we used the ‘vegan::betadisper’ function (Oksanen, 2016).

All R code and data necessary to reproduce our analysis has been archived on Figshare (*link here after acceptance*) and released on GitHub (https://github.com/atredennick/usses_water/releases/v0.1). We also include annotated Stan code in our model description in Appendix 2.

3 RESULTS

Three of our five treatment years fell in years of below average rainfall (Fig. 2A). Thus, those three years represent a lower magnitude of absolute change in precipitation experienced by the treatments. Averaged across treatments, ANPP varied from a minimum of 74.5 g m⁻² in 2014 to a maximum of 237.1 g m⁻² in 2016 (Fig. 2C). ANPP was slightly higher in irrigation plots and slightly lower in drought plots (Fig. 2C), corresponding to estimated soil volumetric water content (VWC) differences among treatments (Fig. 2B). Such differences in soil VWC indicate our treatment infrastructure was successful.

Cumulative March-June soil moisture had a weak positive effect on ANPP (mean of $\beta_1 = 0.41$; Fig. 3), but the effect is associated with high uncertainty, with credible intervals that overlap zero (80% BCI = -0.08, 0.86; 95% BCI = -0.45, 1.13). Ecosystem functional response was similar among treatments, with treatment level regressions between soil moisture and log(ANPP) having similar intercepts and slopes (Fig. 3A). There was also no evidence that the treatment effects became more important over time because there was no directional trend in the random year effects (Fig. 4).

Community composition was similar among treatments. In no year did community composition among treatments not overlap, and they were equally dispersed in all years (Table 1; Fig. 5). Likewise, community composition was remarkably stable over time, with no evidence of divergence among treatments (Table 1; Fig. 5).

4 DISCUSSION

Ecosystem response to precipitation extremes depends on the physiological responses of constituent species and the rate at which community composition shifts to favor species better able to take advantage of, or cope with, new resource levels (Smith et al., 2009). Previous work has shown that community compositional shifts can be both rapid, on the order of years (Hoover et al., 2014), and slow, on order of decades (Knapp et al., 2012; Wilcox et al., 2016). Thus, a lingering question is how the time scales of ecosystem response and community change differ among ecosystems, which can be answered by manipulating precipitation to reach extreme levels. The results of our five year experiment in a sagebrush steppe conform to the first of our four predictions: neither ecosystem functional response nor community composition changed with chronic alteration in water availability (Fig. 1, top left).

The similarity of ecosystem functional response (Fig. 3) and community composition (Fig. 5) across treatments is surprising because grasslands generally, and sagebrush steppe specifically, are considered water-limited systems. Indeed, we expected ecosystem functional response, community composition, or both to change with precipitation treatment, landing us in any box of Fig. 1 *except* the top left. So, why did our treatments fail to induce ecosystem or community responses? We can think of three reasons; two are limitations of our study, and one is the life history traits of the species in our focal communities. We first discuss the potential limitations of our study, and then discuss the biological explanation.

First, it could be that our manipulations were not large enough to induce a response. That is, maybe a 50% decrease/increase in any given year is not abnormal given our site's historical range of variability (Knapp et al., 2017). We cannot definitively rule out this possibility, but we have reason to believe our manipulations *should* have been large enough. Using the methods described by Lemoine et al. (2016), we calculated the percent reduction and increase of mean growing season precipitation necessary to reach the 1% and 99% extremes of the historical precipitation regime at our site. The 1% quantile of precipitation at our site is 110 mm, a 47% reduction from the mean, and the 99% quantile is 414 mm, a 77% increase from mean growing season precipitation (Appendix 3). Thus, our drought treatment represented extreme precipitation amounts, especially in years where ambient precipitation was below average (Fig. 2A). The irrigation treatment may have been too small, however.

Second, ANPP at our site may be influenced by factors beyond the window of soil moisture we included in our statistical model. For example, temperature can impact ANPP directly (Epstein et al., 1997) and by exacerbating the effects of soil moisture (De Boeck et al., 2011). Measurements of soil moisture likely contain a signal of temperature, through its impact on evaporation and infiltration, but the measurements will not capture the direct effect of temperature on metabolic and physiological processes. Likewise, we did not redistribute snow across our treatments in the winter, and snow melt may spur early spring growth. These statistical issues of missing potentially important covariates could explain the weak and uncertain relationship we observed between soil moisture and ANPP.

Third, the life history traits of the dominant species in our study ecosystem may explain the consistent but weak and uncertain effect of soil moisture on ANPP (Fig. 3). Species that live in variable environments, such as cold deserts, must have strategies to ensure long-term success as conditions vary. One strategy is bet hedging, where species forego short-term gains to reduce the variance of long-term success (Seger, 1987). In other words, species do the same thing every year, with only minimal response to environmental conditions. The dry and variable environment of the sagebrush steppe has likely selected for bet hedging species that can maintain function at low water availability and have weak responses to high water availability. In so doing, the dominant species in our ecosystem avoid "boom and bust" cycles, which corresponds to the weak effect of soil moisture on ANPP (i.e., the credible intervals for β_1 overlapping zero), even at precipitation extremes.

Another strategy to deal with variable environmental conditions is avoidance, which would result in a consistent ecosystem functional response at low soil moisture. The perennial plants in this cold desert ecosystem are tolerant to drought conditions (Bazzaz, 1979; Franks, 2011, A.R. Kleinhesselink, unpublished data). For example, many of the perennial grasses in our focal ecosystem avoid drought stress

301 by growing early in the growing season (Blaisdell, 1958, A.R. Kleinhesselink, personal observation).
302 Likewise, the dominant shrub in our focal ecosystem, *Artemisia tripartita*, has access to water deep in the
303 soil profile thanks to a deep root system (Kulmatiski et al., 2017).

304 In conclusion, our results suggest that five years of $\pm 50\%$ ambient precipitation is not enough
305 to induce a shift in ecosystem functional response in a sagebrush steppe. This is likely because the
306 species in our focal plant community are tolerant of drought conditions and bet hedgers in wet conditions,
307 maintaining relatively consistent responses to interannual variation in precipitation to avoid booms and
308 busts. Longer time series of chronic precipitation alteration may reveal plant community shifts that we did
309 not observe (e.g., Wilcox et al., 2016), in which case species that do not bet hedge may gain prominence
310 and dominate the ecosystem functional response. Our results suggest compositional shifts would have the
311 largest impact at high rainfall because the current community maintained consistent ecosystem functional
312 response at very low water availability.

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324 **7 AUTHOR CONTRIBUTIONS**

- 325 • Andrew T. Tredennick collected data, analyzed the data, wrote the paper, prepared figures and/or
326 tables, reviewed drafts of the paper.
- 327 • Andrew R. Kleinhesselink conceived and designed the experiments, performed the experiments,
328 collected data, analyzed the data, reviewed drafts of the paper.
- 329 • Bret Taylor contributed reagents/materials/analysis tools, reviewed drafts of the paper.
- 330 • Peter B. Adler conceived and designed the experiments, performed the experiments, collected data,
331 analyzed the data, reviewed drafts of the paper.

332 **8 SUPPLEMENTAL INFORMATION**

333 **Appendix 1.** Additional methods and information on estimating aboveground net primary productivity.

334 **Appendix 2.** Details of the hierarchical Bayesian model and Fig. A2-1.

335 **Appendix 3.** Details on analysis of precipitation historical range of variability and Fig. A3-1.

9 TABLES

Table 1. Probability that the difference between treatment-level coefficients is greater or less than zero for each of the precipitation treatments. Probabilities denote $\max(\Pr(\text{difference}) > 0, \Pr(\text{difference}) < 0)$, where the sign of the coefficient was ignored. Thus, probabilities represent the probability of the compared coefficients being different from one another. $\Pr(\text{difference})$ values far from 0 or 1 indicate no support for the difference being greater than zero. Bold coefficients denote $\Pr(\text{difference}) \geq 0.95$ (there are none).

Comparison	Intercept	Soil Moisture Effect
Control - Drought	0.70	0.72
Control - Irrigation	0.57	0.85
Drought - Irrigation	0.65	0.70

Table 2. Results from statistical tests for clustering and dispersion of community composition among precipitation treatments. ‘adonis’ tests whether treatments form unique clusters in multidimensional space; ‘betadisper’ tests whether treatments have similar dispersion. For both tests, P values greater than 0.05 indicate there is no support that the treatments differ.

Year	Test	n	d.f.	F	P
2011	adonis	21	2	1.02	0.42
2011	betadisper	21	2	2.23	0.14
2012	adonis	22	2	1.10	0.34
2012	betadisper	22	2	0.21	0.81
2013	adonis	22	2	1.23	0.14
2013	betadisper	22	2	0.28	0.76
2014	adonis	22	2	0.95	0.54
2014	betadisper	22	2	0.35	0.71
2015	adonis	21	2	1.05	0.40
2015	betadisper	21	2	3.01	0.07
2016	adonis	21	2	1.07	0.33
2016	betadisper	21	2	0.50	0.62

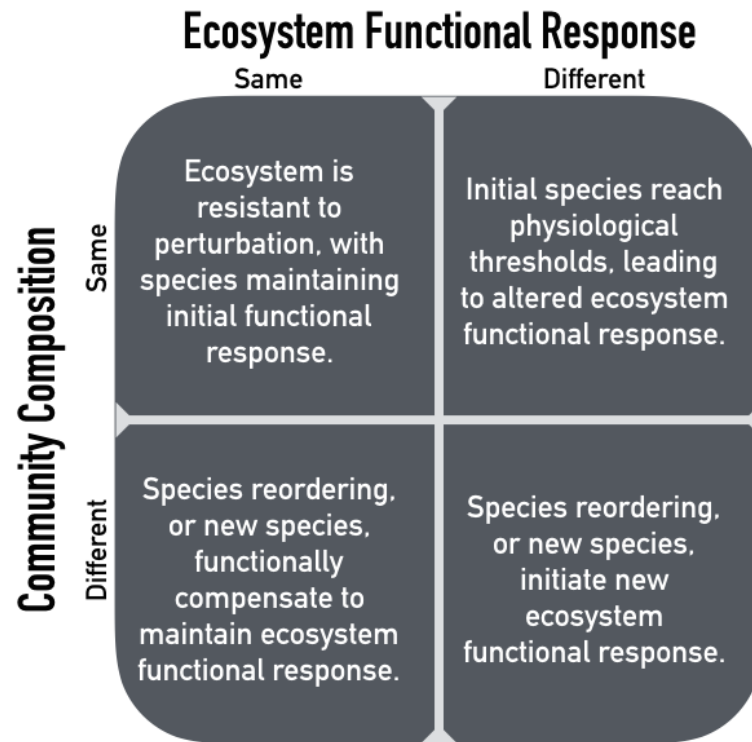


Figure 1. Possible outcomes of chronic resource alteration.

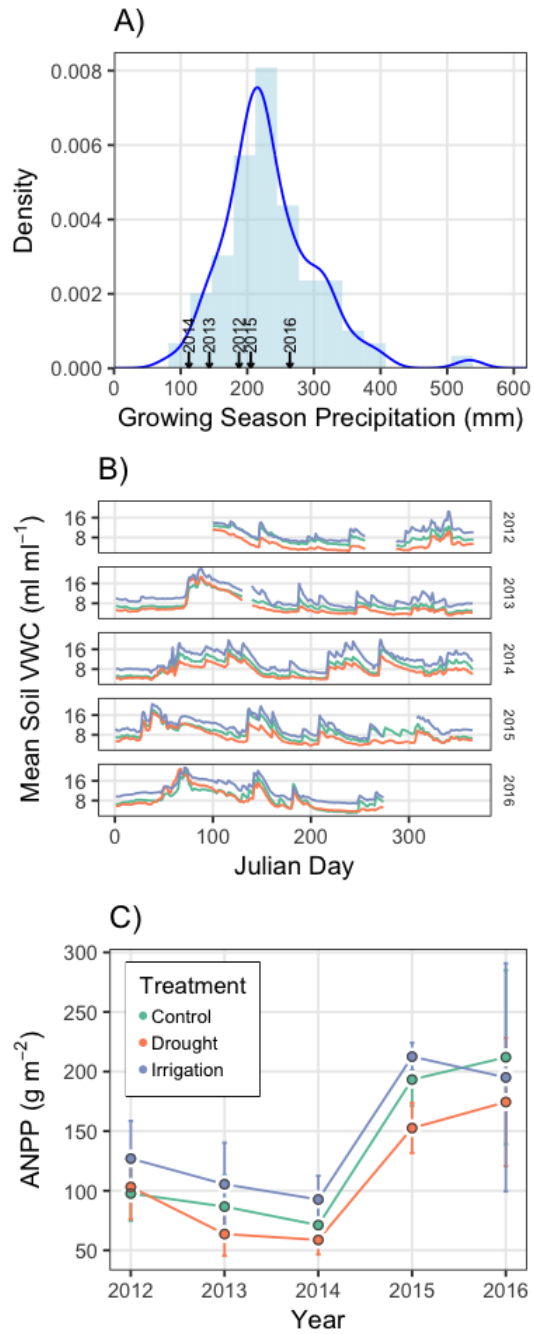


Figure 2. (A) Probability density of historical precipitation from 1926-2016, with the years of the experiment shown with arrows on the x-axis. (B) Observed soil volumetric water content (VWC) over the course of the experiment. (C) Mean (points) ANPP and its standard deviation (error bars) for each year of the experiment.

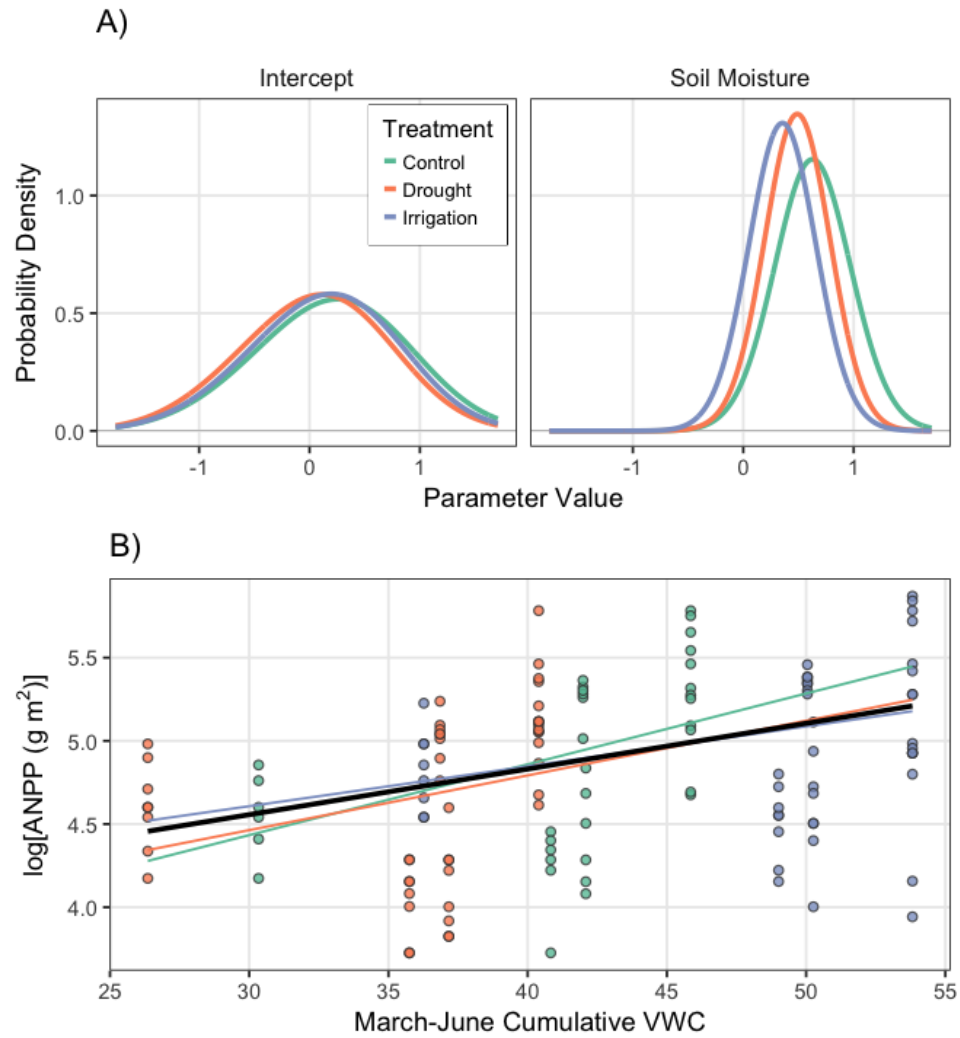


Figure 3. Results from the generalized linear mixed effects model. (A) Posterior distributions of treatment-level parameters ('Intercept' and the effect of 'Soil Moisture'). Kernel bandwidths of posterior densities were adjusted by a factor of 5 for visual clarity. (B) Scatterplot of the data and model estimates shown a solid lines. Model estimates come from treatment level coefficients (colored lines) and the overall mean coefficients (heavy black line). Note that we show log(ANPP) on the y-axis of panel B; this same plot can be seen on the arithmetic scale in supporting material Fig. A2-1.

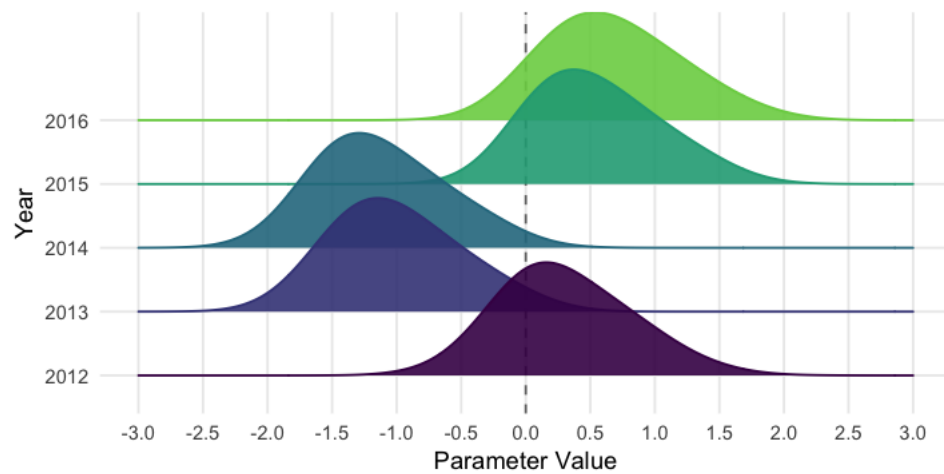


Figure 4. Posterior distributions of random year effects (intercept offsets). Kernel bandwidths of posterior densities were adjusted by a factor of 5 for visual clarity.

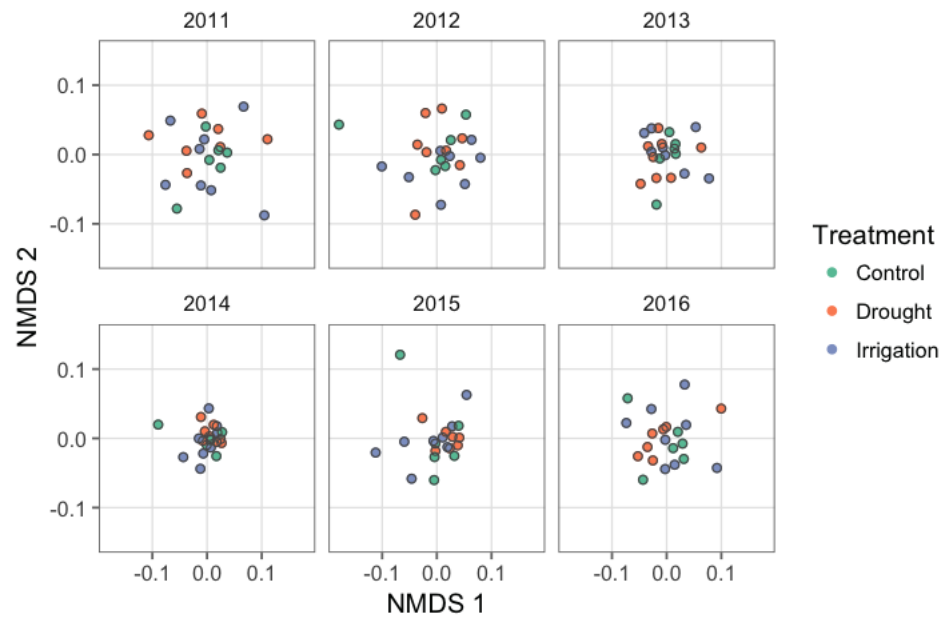


Figure 5. Nonmetric multidimensional scaling scores representing plant communities in each plot, colored by treatment.

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