Effects of climate change and grassland management on plant population dynamics are
mediated by changes in reproductive success

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Abstract

1. Climate change is one of the largest threats to grassland plant species, which can be

modified by land management. Although climate change and land management can

separately and interactively influence plant demography this has been rarely

considered within one experimental set-up.

2. We used a large-scale experiment to quantify the effects of grassland management,

climate change and their joint effect on the demography and population growth rate

of 11 native plant species. We parameterized integral projection models with four

years of demographic data to project the population growth rate. We hypothesized,

plants would perform better in ambient than in the future climate treatment that

creates hotter and drier summer conditions and that plant performance in grazing

vs. mowing would vary across species and depend on their traits.

3. Due to extreme drought events, over half of our study species went quasi extinct. Of

the persistent species, only one supported our expectations, and the rest either had

higher population growth rates in the future climate treatment or showed no

significant difference in population growth between climate treatments. Species

with shorter flowering durations performed better in the mowing treatment while

those with longer flowering durations performed better in the grazing treatment.

The population growth rates of these species were sensitive to changes in vital rates

related to reproduction and recruitment.

4. Our study highlights how extreme climate events, can influence long term

experimental results. Depending on the species we found effects of land

management and climate change on population growth rates but we did not find

strong support for interactive effects among both factors.

5. Experiments combined with measuring plant demographic responses provide a way

to isolate the effects of different drivers on the long-term persistence of species, and

to identify the demographic vital rates that are critical to manage in the future. Our

study suggests that it will become increasingly difficult to maintain species with

preferences for moister soil conditions, that traits such as flowering duration might

predict responses to management, and that vital rates such as reproduction and

recruitment are disproportionately important.

Keywords

Integral projection model, life table response experiment, sheep grazing, mowing, grassland

management, climate change experiment, plant population dynamics, elasticity analysis

Introduction

Grasslands cover approximately 40% of the Earth's surface (Bardgett et al., 2021) and provide

important ecosystem services to society, such as food for livestock and humanity, pollination

and biodiversity (Biurrun et al., 2021; Feurdean et al., 2018). In Central Europe, semi-natural

grasslands were historically managed using low-intensity (i.e. extensive) mowing or grazing,

which prevents their succession into forests (Poschlod et al., 1998; Tälle et al., 2015). These

different types of management affect which types of plants can persist. For example, grazing

favors plants with traits related to higher grazing resistance (e.g. hairiness) or grazing tolerance

(e.g. resprouting capacity) (Bernhardt-Römermann et al., 2011; Evju et al., 2009; Kahmen et al.,

2002). Anthropogenic climate change may modify the effects that management has on the

demography and persistence of plant species (Bardgett et al., 2021; Kariyeva et al., 2012;

Redlich et al., 2022). For example, individuals that currently avoid loss of reproduction from

mowing by flowering before or after mowing events might shift in their phenology in the future

due to climate change, which could lead to mismatches between flowering timing and mowing

events that harm plant reproductive success (Cleland et al., 2006; Gordo & Sanz, 2010).

In central Europe, temperature is projected to increase and it is projected to become wetter in

spring and fall and drier in summer (Döscher et al., 2002; Jacob & Podzun, 1997; Rockel et al.,

2008; Wagner et al., 2013). Drought-tolerant species should be best able to survive the drier

summers expected in the future (Belovsky & Slade, 2021). Species that flower earlier and across

a longer time period might be favored in the future, because they can take best advantage of

the wet spring conditions and because the long flowering duration might buffer their

populations from complete reproductive failure during periods of extreme weather. This leads

to the expectation that drought-tolerant plants with an early flowering start combined with a

long flowering period should be best able to avoid losing reproductive success under future

climate conditions in the presence of grazing and mowing land management types.

In order to understand population dynamics of plant species, demographic studies quantify

vital rates such as survival, growth and reproductive output across the entire life cycle of the

plant and project long-term population growth rate using structured population models

(Caswell, 2000; Crone et al., 2011; Easterling et al., 2000). A common approach to analyzing

plant demographic data are Integral Projection models (IPMs), which consider a relationship

between continuous plant size and vital rates and can also incorporate discrete development

stages (Childs et al., 2003; Jacquemyn et al., 2010; Yule et al., 2013). Retrospective analyses of

IPMs, such as life table response experiments, decompose the contribution of individual vital

rates to the changes in population growth rate between treatments. Vital rates that are more

responsible for the change are those that, either change dramatically between the treatments

and/or those that the population growth rate is sensitive to changes in (Rees & Ellner, 2009).

Many demographic studies have been conducted to assess how the demography of species

responds to climate (Ehrlen, 2019; Ehrlén et al., 2016; Ehrlén & Morris, 2015; Lemmer et al.,

2021; Morris et al., 2020). Most studies focus on natural variation across space and time in

climate conditions and measure plant demographic responses (Compagnoni et al., 2021). Such

observational studies are limited in their ability to assess cause and effect relationships, as

there are often other important factors that co-vary with climate. A few studies have

experimentally assessed how climate affects plant demography (Compagnoni & Adler, 2014;

Lemmer et al., 2021; Lyu & Alexander, 2022; Töpper et al., 2018; Williams et al., 2007).

However, most climate change experiments consider extreme manipulations of temperature

and precipitation, making it hard to predict how plants might respond under realistic future

climate change scenarios (Korell et al., 2020).

We have a unique opportunity to study how a realistic future climate scenario as well as two

different grassland management types (extensive grazing vs. mowing) influence the

demography and population dynamics of plants using the Global Change Experimental Facility

(GCEF) in Germany. Our study builds on the research of Lemmer and colleagues (Lemmer et al.,

2021), which discovered that land management types and climate change interactively affected

the population growth rate of the drought-tolerant grass, *Bromus erectus* L., and that this

interaction was primarily due to higher seedling recruitment of plants under the future climate

and mowing management treatment combination. Lemmer et al. (2020) measured the

demography across a short time period (1transition, two years) and concentrated on a single

species. In this study, we expand the scope to understand the demography and population

responses to climate and land management types across multiple transitions along a longer

time period and multiple plant species. Demographic data was collected over the course of five

years (2018 – 2022) for eleven (Table 1) grassland plant species that represent a range of

different plant statures (tall vs. small), life forms (grasses, herbs) and life histories (biennials vs

perennials). In general, we expected i) the effects of land management types will vary across

species and depend on their traits, ii) plants show higher population growth rates under

ambient compared to future climate conditions and iii) the effects of climate treatments and

land management types on the population growth rate will be interactive.

Methods

Study site

We conducted our research in the Global Change Experimental Facility (GCEF), which was

established in 2013. The experiment is part of the research station Bad Lauchstädt (51°22060 N,

11°50060 E, 118 m a.s.l.). It is designed to simulate a realistic future climate scenario combined

with the manipulation of different land management types, including extensively managed

grasslands. The future climate treatment is based on a mean of 12 regional climate models for

the region (Schädler et al., 2019).

The GCEF consists of 10 main units, five with ambient climate and five with future climate

conditions. Precipitation in the future climate treatment is manipulated seasonally: increases in

precipitation by +10 % compared to the ambient climate in spring and fall and reduced

precipitation by -20 % compared to the ambient climate in summer (for more details see

Schädler et al. 2019). The minimum temperature in the future climate treatment is increased by

about +1.5°C by passive nighttime warming (Schädler et al. 2019).

Each main unit is divided into five subunits that are 16m x 24m in size, each representing a

different land management. Each subunit has 14m x 19.5m available for measurements and the

remainder is left as an edge area, where no experiments or measurements can be undertaken

(Appendix 1, Fig. A1). Our study was established in two of the five land managements, the

extensive meadow (mown one or two times per year) and the extensive pasture (grazed by 20

sheep for 24 hours twice to three times per year). These two land management types were

initially sown in 2013 with 56 plant species native to Germany (Schädler et al., 2019). This

design leads to the following treatment combinations in our experiment: (1) ambient grazing,

(2) ambient mowing, (3) future grazing and (4) future mowing.

Study species

In 2018, we chose eleven different species (Table 1) for our demographic study based on the

criteria that they were abundant enough for demographic data collection (minimum number of

individuals in 2018 was 50 per treatment combination), and that they were species that are

typically associated with mesic or dry grasslands. The species covered a range of growth forms,

plant families and life histories (biennial vs. perennial). Every plant species in this study is native

to Germany and was sown into the GCEF experiment when it was established.

Table 1 Description of each plant species in the study. Life histories are: p = perennial and iteroparous, b

= biennial, h = pluriennial-hapaxanthic (https://wiki.ufz.de/biolflor/index.jsp). Ellenberg wetness values

(indicator of drought tolerance, (Ellenberg et al., 1991): 3 = species is more common at dry sites, avoids

wet habitats (indicator for dry habitats), 4 = usually more on dry habitats, however able to grow on wet

sites, 5 = prefers intermediate locations, missing on very dry and very wet locations, 6 = prefers semi

wet to wet locations, 7 = prefers wet locations, without waterlogging. Ind = indifferent can grow in dry

and wet conditions. In 2020 we recorded the starting month and ending month of flowering for each

species in the GCEF. Flowering duration is the number of months we found an individual flowering

(difference between the end and start month of flowering). Unit of measurement describes how plant

size was measured for each species: basal area (length and width of an individual was recorded and area

was calculated), number of leaves. The grazing resistance of each species was based on the Biolflor

database's index of grazing resistance: 2 = intolerant to sensitive to grazing, 4 = sensitive to moderately

tolerant to grazing, 5 = moderately tolerant to grazing and 6 = moderately tolerant to well tolerant to

grazing, NA = no information available. The mowing resistance of each species was based on the Biolflor

database's index of mowing resistance: 3 = sensitive to mowing, 4 = sensitive to moderately tolerant to

mowing, 5 = moderately tolerant to mowing, 6 = moderately to well tolerant to mowing, 7 = well

tolerant to mowing.

Species name	Growth form	Plant family	Life history	Ellenberg	Start of flowering	End of flowering	Duration of flowering	Unit of measure	Grazing resistance	Mowing tolerance
Anthoxanthum odoratum L.	Grass	Poaceae	p	ind	April	April	1	Basal	5	7
Bromus erectus Huds.	Grass	Poaceae	p	3	May	June	2	Basal	4	5
Crepis biennis L.	Forb	Asterace	b	6	June	August	3	Number	2	6
Dianthus carthusianorum L.	Forb	Caryoph	p	3	May	October	6	Basal	4	3
Lotus corniculatus L.	Legume	Fabacea	p	4	June	July	2	Basal	4	6
Lychnis flos-cuculi L.	Forb	Caryoph	p	7	NA	NA	NA	Basal	2	4
Medicago falcata (L.) Arcang	Legume	Fabacea	p	3	June	July	2	Basal	2	5
Plantago lanceolata L	Forb	Plantagi	p	ind	May	July	3	Number	6	7
Scabiosa ochroleuca L.	Forb	Dipsacac	p	3	June	October	5	Basal	NA	NA
Tragopogon orientalis L.	Forb	Asterace	h	5	June	August	3	Number	NA	NA
Trifolium pratense L.	Legume	Fabacea	p	5	May	May	1	Basal	4	7

Demographic data collection

In April 2018, we established one transect on all our study subunits resulting in 20 transects (five replicates per treatment combination). Our goal was to find a minimum of 10 individuals per transect, to reach a minimum of 50 individuals per treatment combination for every species. Along each transect, 50 x 50 cm subplots were established. We assigned three of the subplots to a fixed positions along each transect (0– 0.50 m; 3 – 3,50 m; 6-6.50 m). The location of other subplots were chosen along the transect at strategic locations to capture individuals of species that were needed to achieve our target of 50 individuals per species and transect. Following this procedure, most transects had eight subplots. This leads to a different overall subplot number for each species, with rarer species having more subplots than common species. Each subplot was marked with a land marking (pipe with red plastic top) in the bottom left and a nail with a white plastic ring the top right corner.

In April 2018, we measured the size of each individual target species in each subplot using either basal area or number of leaves (Table 1). Basal area (cm^2) measured as the product of the horizontal width and length at the base of each individual. Length was measured as the longest lateral dimension of each plant, and width was measured perpendicular to the length. For rosette species (e.g. *Plantago lanceolata, Tragopogon orientalis*), we used the number of leaves as proxy for their size. We recorded the location of each individual within the 50 x 50 cm subplot, using an XY-coordinate system so that individuals could be followed over the years. In rare cases if we spotted individuals of our rarer species just outside of our subplot, we would measure those individuals and track their position relative to the subplot (one of the coordinates would exceed 50 cm).

From 2019 until 2022, we went to the field five times a year. In April, we re-located individuals

from the previous year and measured their size, identified new individuals and measured their

size, and counted the number of new seedlings (for details on how we identified seedlings see

Appendix 3, Table A2). Depending on the phenology of the species and the management events

(see Table 1, Table A1), we counted flowers and seed heads of all reproductive individuals in

each transect. Further we collected two seed heads per subunit and species from individuals

that were outside of our transects but within the same subunits. Number of seeds in each seed

head were counted in the lab. In late autumn (October/November), we counted the number of

seedlings in our subplots, as seedlings of these species can germinate in spring or in fall.

From 2019 onwards, we decided to mark individuals of the very abundant species *Bromus*

erectus with ID's as that helped separating and finding individuals. To mark the individuals, we

used rings made of wire with a tag with a running number on it and put it around the base of

the individual. We left enough space for the individual to grow. Special care was taken to not

harm any of the vegetation during that process.

The multiple droughts Germany experienced during several of the years of this study prevented

more frequent mowing and grazing than would otherwise be incorporated into the treatment

plan (Schädler et al., 2019). During the time of our experiment, the extensive meadows were

mown once per year in early summer time from 2018 – 2020 and two times (early and late

summer) in 2021. The extensively grazed subplots were grazed twice per year in spring and

early summer from 2018-2020, and three times in 2021 (spring, early and late summer). For a

detailed look at the management timing, see Appendix 2, Table A1.

For information on traits for our species, we used Ellenberg values (Ellenberg et al., 1991), the

Bioflor database (https://wiki.ufz.de/biolflor/index.jsp) and our own measurements of

phenology. For drought tolerance, we used categorical Ellenberg values (Ellenberg et al., 1991,

Table 1). For grazing and mowing resistance, we used categorical values from the Bioflor

database (Table 1). In 2020, we collected phenology data on our target species in the GCEF

between April and December within the central 3 x 3 m vegetation survey plot of each subunit

where no experiments or other disturbances (besides the management events) occur. This

provides information on the month in which flowering began and the month it ended, taking all

individuals inside the central vegetation plot into account.

Quasi extinction

Some of the species declined so significantly over the course of our demographic study, that we

considered them quasi extinct and no longer continued to collect demographic data. We

defined quasi extinction from a treatment combination as fewer than 25 individuals total and

fewer than 10 flowering individuals, thus those species treated the species as quasi extinct.

Life cycle stages and vital rates

For all species, we modeled a year-to-year life cycle (April to April) with two stage classes. A

discrete class for seedlings and a continuous class for plants (Fig. 1). Most of our species do not

form long lasting seed banks, and thus we do not have a seed bank class. *Plantago lanceolata*

does have a long-lived seed bank (Chen, 2018), but most of the seeds of this species germinate

immediately and therefore demographic studies on this species do not typically incorporate the

seed bank stage (e.g., www.plantpopnet.com).

Fig. 1 Life cycle diagram of the study species. The diagram shows every vital rate incorporated in the

integral projection model (IPM) and the abbreviation used.

We modeled each continuous state of the IPM as a function of the natural logarithm of the size

of an individual. We used generalized linear models to fit all continuous variables. Survival is a

function of whether or not an individual (i) was still alive at t1, dependent on its log

transformed size (z) at t0 (Eq. 1), modeled as a Bernoulli process with probability of survival S_{t1}

(Eq. 2).

[1] $S_{i,t1} \sim Bernoulli\left(\hat{S}_{t1}\right)$

[2] $logit\left(\hat{S}_{t1}\right) = \alpha + \beta\, log_e(z)$

In this function α is the intercept and β the slope of the curve.

Growth of plants $G_{i,\,t1}$ is described as the normally distributed change in z and log transformed

size at t1 (z') of a surviving individual plant (i) (Eq. 3). It was modeled as a linear function of z' of

surviving individuals depending on z, where the intercept is defined as α^G, the slope as β^G (Eq.

4) and the standard deviation as σ_G (Eq. 3).

[3] $G_{i,t1} \sim Normal\left(\hat{G}_{t1}, \sigma_G\right)$

[4] $\hat{G}_{i,t1} = \alpha^G + \beta^G \, log_e(z)$

Similar to survival, we modeled reproduction probability ($P_{i, t0}$) as Bernoulli process (Eq. 5). Here

we tested how the probability an individual i reproduces at time t0 depends on z. The intercept

is defined as α_t^P and the slope as β^P using a logit link function (Eq. 6)

[5] $P_{i,to} \sim Bernoulli\left(\hat{P}_t\right)$

[6] $logit\left(\hat{P}_{to}\right) = \alpha_t^P + \beta^P \, log_e(z).$

For every reproductive plant, we calculated the number of seeds it produced (Eq. 7). The

number of seeds produced by an individual I at t0 was a function of z, modeled as a Poisson

distribution. The intercept was defined as α_t^F and slope β^F (Eq. 8)

[7] $F_{i,t0} \sim Poisson\left(\hat{F}_{t0}\right)$

[8] $\hat{F}_t = \alpha_t^F + \beta^F \, log_e(z).$

Recruitment, the emerged seedlings per total number of seeds, was separated into fall - ($\theta_{f, j, t0}$)

and spring recruitment ($\theta_{s, j, t0}$). To calculate recruitment, we summed the total number of seeds

per subplot ($S_{j,t}$, Eq. 9). Where $S_{i,t1}$ is the total number of seeds produced by one individual (n) in

a subplot at time t1.

[9] $S_{j,t} = \sum_1^n S_{i,t1}.$

We then divided the result of Eq. 9 by the number of seedlings found in that subplot in fall t0

($Rf_{j,\ to}$, Eq. 10) and spring t1 ($Rs_{j,t1}$, Eq. 11).

[10] $\theta_{f,j,t} = \hat{R}f_{j,to} / S_{j,t}.$

[11] $\theta_{s,j,t+1} = \hat{R}s_{j,t1} / S_{j,t}.$

Establishment (B_j) is the proportion of seedlings that survive to become plants from t0 to t1 in

each subplot j. We calculated the sum of seedlings in spring and fall t0 ($Rsum_{j,\ t0}$) of a subplot

and divided that number by the number of new individuals at t1 ($Ni_{j,\ t1}$) in the same subplot.

[12] $B_j = Ni_{j,t1} / Rsum_{j,t0}$

We calculated the log size distribution of new plants η which is defined as the normally

distributed size of individuals that entered the continuous plant stage in year t1 (Eq. 12). We

used the mean ($log_e(\hat{η}_{t1})$) and standard deviation ($\sigma_η$) of this size in our final parametrization

of the IPM.

[13] $log_e(η_{i,t1}) \sim Normal\left(log_e(\hat{η}_{t1}), \sigma_η\right)$

Effects of treatments on vital rates

For each vital rate (survival, growth, reproduction probability, number of seeds, recruitment,

establishment, size distribution of new plants), we generated three independent pipelines. The

first pipeline differentiated between the climate treatment and thus we calculated the

parameters for ambient and future climate, respectively (Table 2). The second pipeline

differentiated between land management types (mowing and grazing). In the third pipeline, we

differentiated between all four treatment combinations to also account for possible

interactions among treatments. In each of those models, we dropped year, to make the models

more robust (but see Appendix 5 for the "yearly" models).

Table 1. Vital rates included in integral projection model (IPM), its abbreviation, the parameters used

and the according distribution, and which life table response experiment LTRE stage class it was part of.

Vital rates	Abbreviation	Life-cycle stage	Parameters	Distribution	LTRE stage
Plant survival	S	Survival	Intercept (Int), Slope	Bernoulli	Survival
Plant growth	G	Growth	Intercept (Int), Slope, SD	Normal	Growth
Reproduction probability	P	Reproduction	Intercept (Int), Slope	Bernoulli	Reproduction
Number of seeds per reproductive plant	F	Reproduction	Intercept (Int), Slope	Poisson	Reproduction
Fall recruitment	θf	Recruitment	Mean		Recruitment
Spring recruitment	θs	Recruitment	Mean		Recruitment
Seedling establishment	B	Establishment	Mean		Establishment
Size distribution of new plants	η	Establishment	Mean, SD		Establishment

Integral projection model (IPM)

For each species and pipeline, we created an IPM to calculate the influence of climate and

management on the long-term population growth (λ). In every IPM, we calculate all possible

transitions from z to z' at. The change in number of plants from one year to the other is described

by:

[14] $$n(z', t1) = M(t0)B\eta(z') + \int_{L}^{U} S(z)G(z', z) + P(z)F(z)\theta_f B\eta(z')n(z, t0)dz$$

Where the vector n(z′, t1) is the number of plants at size z′ at time t1. In this function, the first

term represents the recruitment of spring seedlings entering the adult plant class. This is based

on the number of spring seedlings at t0, M(t0), the seedling establishment, B, and the size

distribution of new plants η(z′). The second kernel (or surface) describes the growth and survival

of plants from t0, n(z, t0) to t1, n(z′, t1). It is defined as an integral with in the upper limit U, which

is the biggest size observed and the lowest size L, observed for the species in our study. The

integrals were evaluated across 200 equally spaced size bins using the midpoint rules as a 200 x

200 matrix.

The recruitment of spring seedlings from one year to the next is described by:

[15] $M(t + 1) = \int_L^U P(z)F(z)\theta_s dz.$

As we did not distinguish the new plants we see in April as those that were seedlings from fall vs.

spring, we use the same establishment estimate for seedlings (B) twice in the model.

Effects of treatments on population growth rate

To determine the effect of the climate treatment on λ, we calculated λ for the ambient and

future treatments and then calculated an effect size by subtracting λ ambient from λ future.

Positive values show a species´ preference for ambient climate conditions whereas negative

values show a species´ preference for future climate conditions. Likewise, for land management

types, we subtracted λ grazed from λ mowing to calculate an effect size. Here, positive values

311 show a species preference for the grazing treatment whereas negative values show a species

312 favors the mowing treatment.

313 To test whether traits influenced the effect size of land management types on λ, we ran simple

314 linear models using the effect size as response variable and the following traits as explanatory

315 variable: flowering start month, duration of flowering, grazing resistance and mowing

316 resistance. Because flowering time, start and end could differ between treatments due to the

317 effect of the land management type we decided to not calculate a mean across those.

318 To quantify uncertainty in estimates of λ, we bootstrapped our data a thousand times and

319 calculated the λ each time. To test for significant differences in λ between treatments and

320 treatment combinations, we used permutation (randomization) tests (N = 1000 permutations).

321 For each treatment combination, we calculated sensitivities and elasticities of λ to kernel

322 element perturbations.

324 **Life table response experiment (LTRE)**

325 In order to decompose the influences of vital rates on the difference in λ, across treatment

326 combinations, we conducted Life Table Response Experiments (LTRE) for each pairwise

327 treatment combination. The difference in λ, $\Delta\lambda$, is:

328 [17]   $\Delta\lambda = \lambda^{treatment\ combination\ 1} - \lambda^{treatment\ combination\ 2}$

329 The contribution of each vital rate to $\Delta\lambda$, $\widetilde{\delta}^{\,\alpha i}$, is:

330 [18]   $\widetilde{\delta}^{\,\alpha i} = \sum_{i}^{14}(a_i^{treatment\ combination\ 1} - a_i^{treatment\ combination\ 2}) \frac{\partial\lambda}{\partial\alpha_i}$

Here α_i is one of the fourteen vital rates that are included in the IPMs, while the sensitivity of λ

to each vital rate is defined as $\frac{\partial \lambda}{\partial \alpha_i}$. Vital rates will have a strong contribution to the observed

difference in λ between treatment combinations if the vital rates change strongly in magnitude

between the two treatment combinations and/or if λ is sensitive to changes in that vital rate. To

facilitate the understandability, we combined the vital rates with respect to six demographic

processes: survival, growth, reproduction, recruitment, establishment (Table 2). After this we

scaled $\widetilde{\delta}^{\,\alpha_i}$ to 1 (to display proportional influence).

Analysis

All analysis were carried out in R (R Core Team, 2022). Figures were made using the ggplot2

package (Wickham, 2016). The integral projection model was run and calculated using the ipmr

packages (Levin et al., 2021). Other packages used were: openxlsx (Schauberger & Walker,

2021) to load the xlsx files into R, dplyr (Wickham et al., 2022) for structuring data and piping

and the package parallel (R Core Team, 2022) to decrease the running time of R scripts. The R

code can be found at GitHub:

Martin19910130/Effect_of_climate_and_land_use_on_population_growth_rate

and the data is supplied at figshare:

https://figshare.com/account/home#/projects/157925.

Results

Quasi extinction

The following six species went quasi extinct over the course of the experiment: *Anthoxanthum*

odoratum, Crepis biennis, *Lotus corniculatus*, *Lychnis flos-cuculi* and *Trifolium pratense*. While

there is too small of a sample size (N=11 species) for statistical analysis, it is notable that the

species that went quasi extinct were also the ones that were less drought tolerant based on

their Ellenberg values (Table 1) and that the first few years of our study were extraordinarily dry

for our region (Boergens et al., 2020).

Management effects on population growth rate

Tragopogon orientalis had significantly higher λ in mowing compared to grazing management

(permutation test p = 0.042, Fig. 2E), whereas all other species did not show a significant main

effect of management on λ (Fig. 2A-D).

Fig. 2 Log population growth rate (λ) for each of the five study species (A-E) considering the two

different land management types (grazing and mowing). Displayed is the mean $\pm$ one standard

deviation.

Traits related to grazing resistance and mowing resistance did not explain the effect size of land

management type on λ (Appendix 4, Fig. A2). However, flowering duration was positively

related to the effect size of land management type on λ; species with lower flowering duration

(i.e. *Plantago lanceolata, Bromus erectus, Tragopogon orientalis*) performed better in the

mowing treatment while those with longer flowering duration (*Dianthus carthusianorum,*

Scabiosa ochroleuca) performed better in the grazing treatment (t = 6.087, p < 0.001, r^2 = 0.8;

Fig. 3A). The start of flowering was a poor predictor of management preferences (Fig. 3B, t = -

0.206, p = 0.842, r^2 = -0.1191).

Fig. 3 Relationship between the effect size of management on population growth rate (λ grazing − λ

mowing) plotted against A) The mean duration of flowering for each species and B) the mean beginning

of flowering for each species. The line indicates the trend from the linear model. Symbols show the

management treatment in which phenology was measured.

Climate effects on population growth rate

B. erectus had a significantly higher λ in future compared to ambient climate (permutation test,

p = 0.023, Fig. 4A), and the other four species did not show significant differences in λ with

regard to the climate treatment (Fig. 4B-E).

Fig. 4 Bootstrapped logged λ plotted against the two climate treatments (ambient and future) for the

five different study species (A-E). Displayed is the mean logged λ of the 1000 repetitions, with the

standard deviation.

Effect of climate and land management type on population growth rate

In *S. ochroleuca,* λ was significantly higher in future compared to ambient climate in the mowing land management type (future mowing > ambient mowing; permutation test, p = 0.011) but not in grazing land management (permutation test, p > 0.05, Fig. 5D). The λ of *T. orientalis* was significantly higher in ambient compared to future climate only in the grazing management treatment (ambient grazing > future grazing; permutation test, p = 0.006) and λ was higher in mowing compared to grazing management only in future climate (permutation test, p = 0.026, Fig. 5E). For all other species (*B. erectus*, *D. carthusianorum* and *P. lanceolata*) the lines in the interaction plots crossed but they were not significantly different from each other due to the large variability around the mean (Fig. 5A-C).

Fig. 5 Effect of climate (ambient and future) and land management type (grazing and mowing) on population growth rate (λ) of the five study species (A-E). Displayed is the mean and the standard deviation.

Sensitivity

Across all species and treatments, λ was most sensitive to changes in reproduction parameters (Fig. 6). While the magnitude differed between treatments and species (e.g. *S. ochroleuca* highest in grazing and future *D. carthusianorum* highest in ambient).

Fig. 6 Sensitivity of pairwise treatment combinations for every vital rate included in the IPM. Each

species is color-coded.

Life Table Response Experiment

The higher λ of *B. erectus* in future compared to ambient climate was primarily due to increases

in reproduction and recruitment of this species under future climate (Fig. 7C and 7D). The

higher λ of *S. ochroleuca* in future compared to ambient climate in the mowing management

treatment was primarily due to increases in the reproduction of this species in the future

mowing treatment (Fig. 7D). The higher λ of *T. orientalis* in ambient compared to future climate

in the grazing management treatment was primarily due to increases in plant survivorship (Fig.

7C), whereas the higher λ in this species in mowing compared to grazing in future climate

treatment was due to higher survival and reproduction (Fig. 7B).

Fig. 7 Results for the Life Table Response Experiment (LTRE). Each species is displayed in a different

color. A) Ambient climate treatment and the effect of the two land management types on the

parameters. B) Future climate treatment and the effect of the two land management types on the

parameters. C) Grazing land management type and the effect of climate on the parameters. D) Mowing

land management type and the effect of climate on the life stages. For a better overview, we summed

the parameters according to their corresponding live stage.

Discussion

Our study aimed to examine whether plant species interactively respond to climate and land

management types in their demography and population dynamics. However, in the early years

of our study, six of our eleven focal species went quasi extinct from the experiment, likely due

to the extreme drought conditions that occurred in our region in 2018 and 2019. This is

becoming an increasing phenomenon in experimental global change research—even the

control treatments experience extreme climate events because these are now more common in

our rapidly changing world (IPCC, 2014; Jentsch et al., 2007; Orlowsky & Seneviratne, 2012;

Tebaldi et al., 2006). Of the remaining five species, we found that flowering duration is an

important trait that predicts responses to land management types, with shorter duration

species having higher λ in mowing management and longer duration species having higher λ in

grazing management. In contrast to our expectation that plants would perform better in

ambient compared to future climate conditions, we found that two species (*B. erectus* and *S.*

ochroleuca) have higher λ in the future climate treatment and only *T. orientalis* shows the

expected higher λ in ambient climate. We suggest this is likely due to the high drought

tolerance of *B. erectus* and *S. ochroleuca*. We did not find strong support that the effects of

climate and land management types were interactive for our study species, and it is unclear if

this is because these interactive effects do not occur or because we have low power to detect

them.

We expected that plants species in the study region (Central Germany) would perform best in

ambient climate, as they would have difficulty dealing with the dry conditions expected in the

summer under future climate. However, our results do not support this hypothesis. The first

year of our demographic study, 2018, coincided with the driest summer for Germany since start

of recording (Schuldt et al., 2020). This drought had dramatic and long-lasting effects on the

plant communities across the entire GCEF experiment. Many of the species that went extinct

were those that are indicative of moist soil conditions. Four of the five species that remained

are indicative of drier habitat conditions and commonly occur in semi-dry or dry meadows (*B.*

erectus, *D. carthusianorum* and *S. ochroleuca*) or are known to be highly generalized with

regards to the moisture of the habitat (*P. lanceolata*). *B. erectus* and *S. ochroleuca* were also

the two species that performed better in future experimental climate conditions. This result

matches previous experimental studies (Lemmer et al., 2021; Moser et al., 2011; Poniatowski et

al., 2018) that show high performance of *B. erectus* in hot and dry summer conditions. Our LTRE

results revealed that the higher λ of *B. erectus* in future climate was due primarily to the higher

reproduction and recruitment in this treatment (see also Lemmer et al. 2021). The earlier

flowering of *B. erectus* compared to other species could explain its relative advantage in future

climate conditions; *B. erectus* is able to take advantage of the additional water resources in

spring and reproduce before the hot and dry summer conditions in the future climate

treatment.

Only one of our remaining species, *T. orientalis,* is indicative of moist to moderately dry soil

conditions (Sebald et al., 1990; Ellenberg et al., 1991; Bomble, 2013) and managed to persist in

our study. This species did indeed show the expected higher performance in the ambient

compared to future climate treatment. We suggest that we may have found more support for

our hypothesis if we had been able to study the demography of more typical species from

moister grassland communities as we initially planned. Our study highlights the importance of

historical legacies in shaping experimental results and support projections that communities

may shift towards more drought tolerant species under future climate conditions (Belovsky &

Slade, 2021; Craine et al., 2013).

Surprisingly, most of our study species did not show strong differences in their population

growth across land management types. The only exception was the biennial species, *T.*

orientalis, which performed significantly better in the mowing compared to grazing

management treatment, primarily due to higher survival and reproduction in the mowing

treatment. The vegetative parts of *T. orientalis* are relatively soft and easy for sheep to chew

(personal observation) and the large flower heads may attract sheep. We also observed that

the flower buds were eaten by other wild animals, such as rabbits (personal observation). This

species adds to several studies that showed that grazing has a negative effect on plant

population dynamics (Hansen & Wilson, 2006; Jacquemyn et al., 2012; van der Meer et al., 2014). For

example, in *Primula veris*, grazing was found to decrease population growth rate, primarily

through its negative effect on reproduction (Jacquemyn et al., 2012).

Our results show that the effect of management on λ depended on flowering duration. Plants

that flowered a relatively short duration showed higher λ in mowing (e.g. *T. orientalis*), whereas

those flowering across a longer duration showed higher λ in the grazing management. This

could be because plant populations that flower across a long time period are buffered from

losing all of their reproductive success from disturbances that occur in a short time interval,

such as grazing. However, we have to be cautious when interpreting these results as in our

study years the management was mostly applied in a lower frequency than usual; most years

had one mowing and two grazing events. This was because mowing and grazing were not

possible when there was too little biomass regrowth, which occurred during the dry study

years. Other studies have also shown that low frequency (once a year) and late season mowing

benefits the reproduction of plants (Jantunen et al., 2007; Johansen et al., 2019). Across all of

our study species, we found that the population growth rates were most sensitive to

reproduction. Thus, it may be a general finding that management treatments should be timed

to interfere as little as possible with peak flowering and fruiting (Nakahama et al., 2016; Reisch &

Poschlod, 2009).

We expected that interaction effects of treatments on λ would be common in our study and

that each species would have a treatment combination that was most optimal for λ. However,

we had little power to test for these interactions due to the high variation across individuals in

the sensitive vital rates of seed production and recruitment. For example, the high variation in λ

of *S. ochroleuca* in the ambient-grazing treatment combination is likely due to very high seed

production (>2000 seeds) of three individuals in this treatment combination (Appendix 6).

Similarly, the high variation in λ in the ambient treatment for *D. carthusianorum* is likely due to

high seed production of a few individuals in the ambient mowing treatment (Appendix 6) and

high recruitment of new spring seedlings in a few plots in the ambient grazing treatment

(Appendix 7). Finally, while *P. lanceolata* appears overall have higher λ in ambient compared to

future climate treatments, high variance in the future-mowing treatment combination, which is

likely due to two reproductive individuals with outlier high seed production, may prevent the

detection of an effect.

In demographic studies of both plants and animals, reproductive output is known to be skewed

across individuals, with only a few individuals producing many offspring (van Daalen & Caswell,

2020). The ability to distinguish deterministic treatments effects from stochasticity among

individuals would require much higher sample sizes than is available in our GCEF experiment.

While there are many advantages of conducting multi-species demographic studies like this one

and of conducting demographic research in a controlled experimental setting, one distinct

disadvantage is the effort and sample size required to adequately detect small changes in

sensitive vital rates (McMahon et al., 2019).

It was surprising to find in our LTRE that changes in reproduction and recruitment often

contributed most to the observed responses of λ to management and climate. Usually in longer

lived species, λ is highly sensitive to changes in survival (Weppler, Stoll and Stöcklin, 2006; Kuss

et al., 2008) and thus even small changes in survival across treatments should contribute

significantly to changes in λ. In our case, we find that λ is highly sensitive to recruitment. This

might be due to the relatively high λ of our perennial species; species with high λ are often

more sensitive to reproduction and recruitment than to survivorship (Franco & Silvertown, 2004;

Ramula et al., 2008). Further, many of our species were able to maintain their survival across

treatments, but had high changes in their reproduction and recruitment.

Conclusions

Our study is a first step towards answering important yet unanswered question on how climate

change affects population dynamics of grassland species and how this effect is modified by

different land management types (Ehrlen, 2019). Our results are optimistic for our four more

drought-tolerant species characteristic for dry or semi-dry grasslands, showing persistent or

growing populations in both management types (grazing vs. mowing) and under future climate

conditions as well as after natural extreme drought events. However, less drought-tolerant

species either went quasi extinct early in the study or were projected to decline in the future,

indicating that populations cannot persist in the presence of future climate conditions. Our

results suggest that it will be more difficult to create and maintain especially mesophilic

grasslands composed of species with preference for moister soil conditions in the future.

Furthermore, we suggest that management in future climate should carefully be timed outside

of peak flowering and fruiting, so that as many species as possible can reproduce and persist in

managed grasslands.

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Conflict of interest

No Author has a conflict of interest

Data availability statement

Data (Size data and reproduction data, as well as number of seed data):

https://figshare.com/account/home#/projects/157925

Author contributions

The study was designed by LK (leading) and MA and TMK (supporting). Fieldwork was designed

and carried out by LK and MA (both leading). The data for the phenology of plants was collected

by CP (leading). Analysis was carried out by MA (leading) with support of the other co-authors.

The manuscript was written by MA (leading) and TMK and LK (supporting). All authors

commented and improved the manuscript.

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Appendix 1

Fig. A1 Layout of every GCEF plot. 1 indicates where the transect was situated at each GCEF subunit.

Displayed is the overall size of the GCEF subunits and the inner subunit where experiments take place.

Appendix 2

Table A1 Given are the dates of the management events. Sheep graze one plot for 24 hours and thus the

management took place over the course of 10 days. The sheep were let on one plot in the morning of

the day and the next day they moved to the next plot. Mowing was done in one day.

Management event	Year	Date
Grazing	2018	30.04 - 11.05
Grazing	2018	11.06 - 22.06
Mowing	2018	11.06
Grazing	2019	6.05 - 17.05
Grazing	2019	10.06 - 21.06
Mowing	2019	11.06
Grazing	2020	11.05 - 22.5
Grazing	2020	22.06 - 03.07
Mowing	2020	08.06
Grazing	2021	10.05 - 21.05
Grazing	2021	14.06 - 25.06
Mowing	2021	14.06
Mowing	2021	06.09.2023
Grazing	2021	06.09 - 17.09

834 **Appendix 3**

835 **Table A2** Information on how we identified seedlings. Mostly that stage class was size dependent and if

836 an individual plant didn't grow to a certain size, it was identified as seedling. Some *Bromus erectus*

837 individuals flowered after they developed 4 leafs but still were smaller then 0.25 cm^2 which is why we

838 classified every individual with three or more leafs as adult.

Species	Maximum size as a seedling
Anthoxanthum odoratum	< 4 leafs
Bromus erectus	1 ramet with maximum 3 leafs
Crepis bunnies	1 ramet with maximum 3 leafs
Dianthus carthusianorum	Size of rosette <0.25cm^2
Lotus corniculatus	Size of rosette <0.25cm^2
Lychnis flos-cuculi	Didn't find seedling
Medicago falcata	Size of rosette <0.25cm^2
Plantago lanceolata	1 ramet with maximum 3 leafs
Scabiosa ochroleuca	1 ramet with maximum 2 leafs
Tragopogon orientalis	Maximal 3 leafs
Trifolium pratense	1 individual with maximum 2 leafs

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847   **Appendix 4**

848   **Fig. A2** Grazing and mowing resistance plotted against the effect size of λ. Negative values indicate that

849   the species has a higher λ in the mowing treatment. Panel A.) shows the effect of grazing resistance on

850   the effect size and panel B.) shows the effect of mowing resistance on the effect size.

Appendix 5

Fig. A3 Bootstrapped λ of each species over the study years for the management treatment (mowing,

grazing). Displayed is the mean λ and the standard deviation. Each year on the x axis stands for a

transition: 2018 = 2018–2019, 2019 = 2019–2020, 2020 = 2020–2021, 2021 = 2021–2022. For the

transition 2021 we were not able to calculate a lambda for *T. orientalis* in the grazing land management

treatment due to lack of data points.

Fig. A4 Bootstrapped λ of each species over the study years for the climate treatment (ambient, future).

Displayed is the mean λ and the standard deviation. Each year on the x axis stands for a transition: 2018

= 2018 – 2019, 2019 = 2019 – 2020, 2020 = 2020 – 2021, 2021 = 2021 – 2022. S. ochroleuca in 2019 and

T. orientalis did not have enough data points in order for us to calculate the population growth rate.

881 **Fig. A5** The effect of the treatment combinations (management and climate) over the years of the study

882 on λ. Displayed is the mean λ and the standard deviation. Each year on the x axis stands for a transition:

883 2018 = 2018 − 2019, 2019 = 2019 − 2020, 2020 = 2020 − 2021, 2021 = 2021 − 2022. Missing points are

884 due to lack of data points to calculate the population growth rate.

    Appendix 6

892   **Fig. A6** Number of seeds plotted against the size of each reproductive individual for the species (A)

893   *Bromus erectus,* (B) *Dianthus carthusianorum*, (C) *Plantago lanceolata*, (D) *Scabiosa ochroleuca* and (E)

894   *Tragopogon orientalis*. Different treatments are indicated with different shapes and colors.

 Appendix 7

904   **Fig. A7** Turnover from seeds to seedling in the different treatments for (A) *Bromus erectus*, spring

905   recruitment (B) *Bromus erectus*, fall recruitment (C) *Dianthus carthusianorum*, spring recruitment (D)

906   *Dianthus carthusianorum*, fall recruitment (E) *Plantago lanceolata*, spring recruitment (F) *Plantago*

907   *lanceolata*, fall recruitment (G) *Scabiosa ochroleuca*, spring recruitment (H) *Scabiosa ochroleuca*, fall

908   recruitment (I) *Tragopogon orientalis*, spring recruitment (J) *Tragopogon orientalis*, fall recruitment.

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