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# Density feedbacks mediate effects of environmental change on population dynamics of a semi-desert rodent

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## Abstract

- 1. Population dynamics are the result of an interplay between extrinsic and intrinsic environmental drivers. Predicting the effects of environmental change on wildlife populations therefore requires a thorough understanding of the mechanisms through which different environmental drivers interact to generate changes in population size and structure.
- 2. In this study, we disentangled the roles of temperature, food availability, and population density in shaping short- and long-term population dynamics of the African striped mouse, a small rodent inhabiting a semi-desert with high intra- and inter-annual variation in environmental conditions.
- 3. We parameterized a female-only stage-structured matrix population model with vital rates depending on temperature, food availability, and population density, using monthly mark-recapture data from 1609 mice trapped over 9 years (2005-2014). We then applied perturbation analyses to determine relative strengths and demographic pathways of these drivers in affecting population dynamics. Furthermore, we used stochastic population projections to gain insights into how three different climate change scenarios might affect size, structure, and persistence of this population.
- 4. We identified food availability, acting through reproduction, as the main driver of changes in both short- and long-term population dynamics. This mechanism was mediated by strong density feedbacks, which stabilized the population after high peaks and allowed it to recover from detrimental crashes. Density dependence thus buffered the population against environmental change,

and even adverse climate change scenarios were predicted to have little effect on population persistence (extinction risk over 100 years < 5%) despite leading to overall lower abundances.

5. Explicitly linking environment-demography relationships to population dynamics allowed us to accurately capture past population dynamics. It further enabled establishing the roles and relative importances of extrinsic and intrinsic environmental drivers, and we conclude that doing this is essential when investigating impacts of climate change on wildlife populations.

## Keywords

environmental drivers, environmental stochasticity, extrinsic, intrinsic, LTRE, matrix model, perturbation analysis, population dynamics, rodent, vital rate

## <sup>1</sup> Introduction

Contemporary climate change happens at a fast rate, and increases in temperatures 2 and the frequency of extreme events are predicted to continue (IPCC, 2014). Effects 3 of climate change on ecological systems have already become evident (Parmesan, 4 2006) and manifest themselves as alterations in species distributions (Chen et al., 5 2011), shifts in phenology (Charmantier et al., 2008), and changes in abundance 6 (Pounds et al., 2006; Jepsen et al., 2008). These impacts make it more impor-7 tant than ever to understand and predict how climate affects life-history processes 8 and population dynamics of animals and plants both directly and indirectly (e.g. 9

<sup>10</sup> through resource availability)(Williams *et al.*, 2002).

Population dynamics are the result of an interplay between different extrinsic 11 and intrinsic environmental factors (Stenseth et al., 2002; Goswami et al., 2011). 12 Extrinsic factors such as climate and resource availability interact with intrinsic 13 density feedbacks to generate variation in vital rates (survival, reproduction). Vi-14 tal rate variation then translates into changes in population size and structure 15 (Leirs et al., 1997; Lima et al., 1999). The impacts of extrinsic environmental 16 factors and how these are affected by density may differ among seasons (e.g. Gul-17 lett et al., 2014), across species' life cycles (Gamelon et al., 2017) and be subject 18 to stochastic variation among years, making population dynamics dependent not 19 only on current, but also on past conditions (Wilmers et al., 2007). Predicting 20 potential impacts of climate change on species and ecosystems therefore requires 21 an in-depth understanding of how deterministic and stochastic variation in cli-22 mate, resource availability, and density feedbacks jointly affect population dynam-23 ics (Benton et al., 2006; Boyce et al., 2006). Such understanding can be gained 24 through quantitative models that explicitly link environmental factors to popula-25 tion dynamics via vital rates (Ehrlén et al., 2016) and the availability of long-term 26 individual-based data necessary to parameterize these models (Clutton-Brock & 27 Sheldon, 2010). 28

Collecting sufficient amounts of long-term individual-based data is easier for species with short generation time, such as rodents. Rodent life histories are also of particular interest because they are highly sensitive to the environment, and consequently show large numerical fluctuations within and among years (Krebs, 2013). These fluctuations are of great ecological, social, and economical interest. They are, for example, linked to ecosystem consequences of the dampening of vole

and lemming population cycles (Ims et al., 2008; Schmidt et al., 2012), stochas-35 tic rodent eruptions with detrimental effects on crop production (Singleton et al., 36 2010), and high risks of disease outbreaks (Gubler *et al.*, 2001). Rodents are also 37 often keystone species in food-webs and can function as ecosystem engineers (Kelt, 38 2011). Furthermore, while studying rodent populations - particularly in the con-39 text of environmental change - is worthwhile in itself, drivers and mechanisms of 40 population dynamics may also be conserved across species. From a broader per-41 spective, we may thus be able to use studies on easily accessible rodent systems 42 to gain valuable insights into general principles of population dynamics in variable 43 environments and to draw inferences about related or ecologically similar species 44 for which long-term individual-based data are not available (Frederiksen et al., 45 2014). 46

Here, we study population dynamics of the African striped mouse (*Rhabdomys* 47 *pumilio*), a small rodent inhabiting a semi-arid ecosystem characterized by con-48 siderable variation in timing and amount of annual rainfall and, consequently, 49 availability of green vegetation (Cowling et al., 1999). Being adapted to such a 50 variable environment, the species displays high degrees of phenotypic plasticity 51 (Raynaud & Schradin, 2008; Nel et al., 2015) and potentially large adaptive ca-52 pacity in the face of environmental change (Rymer *et al.*, 2013). Variation in vital 53 rates of this rodent has previously been linked to temperature, food availability, 54 and population density (Nater et al., 2016a). In this study, we re-analyse these 55 environment-demography relationships and link them to population dynamics in 56 a stage-structured population model. We then subject this model to retrospective 57 perturbation analysis to identify the roles and relative strengths of environmental 58 drivers in generating past population dynamics, and to prospective perturbation 59

analysis to investigate potential future responses of the African striped mouse to
alterations in the stochastic environment including different scenarios for future
climate change.

## 63 Materials and methods

#### 64 STUDY SYSTEM AND DEMOGRAPHIC DATA

The African striped mouse (Rhabdomys pumilio) is a small murid rodent (adult 65 body mass of 30-85 g in the field) native to the dry regions of South Africa (Mal-66 larino et al., 2018). It is diurnal, forages alone but returns to a nest shared with an 67 extended family group at night (Schradin & Pillay, 2004). Group members share 68 one territory and interact amicably with each other, but are highly aggressive to-69 wards striped mice from other groups (Schradin & Pillay, 2004). Striped mice are 70 omnivores, feeding primarily on the leaves of shrubs, small succulents and ephemer-71 als, but have also been observed to eat seeds and insects (Schradin, 2005). Home 72 range sizes vary strongly depending on local population density and seasonal food 73 availability (Schradin, 2006). The main breeding season is in the austral spring. 74 Most striped mice are therefore born between July and November, and can reach 75 sexual maturity after four weeks of age (Schradin & Pillay, 2014). However, they 76 often delay reproduction and remain in their natal territory until the breeding 77 season of the following year and few mice survive to a another breeding season 78 after that (life expectancy rarely exceeds 2 years, Schradin et al. (2012)). 79

The study population of African striped mice is located in the Goegap Nature Reserve in the Succulent Karoo of South Africa (29°41′ S, 18°01′ E; altitude 912 m), a semi-arid winter-rainfall ecosystem with marked vegetation peaks in spring. The

study population has been monitored since 2004 with a monthly capture-mark-83 recapture program (trapping protocol described in detail in Schradin (2006)). For 84 this study, we assigned female striped mice to one of three life stages based on age 85 and reproductive status. Individuals below the age of four weeks were considered 86 immatures, while those that were older than four weeks but had not yet shown 87 signs of reproduction were assumed to be philopatrics (pre-reproductive adults). 88 Striped mice displaying a perforated vagina in month t-1, and/or signs of lactation 89 in month t were considered breeders (reproductive adults) from month t onwards 90 (Figure 1). 91

#### 92 ENVIRONMENTAL DATA

To investigate environmental effects on vital rates and population dynamics of striped mice, we explicitly included ambient temperature, food availability, and adult population density into our analyses.

<sup>96</sup> Monthly mean temperature represents a key seasonal cue, and was calculated <sup>97</sup> by averaging over daily measurements of minimum and maximum temperatures <sup>98</sup> collected at the field station. Absolute minimum and maximum temperatures in <sup>99</sup> each month were highly correlated with monthly mean temperatures (Pearson's r <sup>100</sup> = 0.784 and 0.868 respectively), and were therefore not considered separately in <sup>101</sup> our analyses.

We quantified monthly food availability using estimated abundance of annual succulents and ephemerals eaten by striped mice (34 species, Schradin (2006)). These plants vary seasonally in abundance and are the main driver of both changes in metabolic rates (Rimbach *et al.*, 2018) and reproduction (Nel *et al.*, 2015) of striped mice. We estimated plant abundance based on a vegetation survey within

the study site. Eight monitoring plots (2 x 2 m each) were sampled monthly using 107 a standard protocol (Braun-Blanquet method, Werger (1974)) to determine the 108 amount of ground covered by different species of annual succulents and ephemerals. 109 Monthly food availability was then calculated as the plot-average percentage of 110 ground covered by all plant species. We left out the shrub components of striped 111 mouse diet, Lycium cinereum and Zygophyllum retrofractum, as the former has 112 annual leaves that covary seasonally with the abundance of annual succulents 113 and ephemerals, while the latter is a succulent that varies little within years and 114 represents an "emergency food" for striped mice when nothing else is available 115 (Schradin, 2006). We also did not consider rainfall directly as a covariate as studies 116 have shown that water is not a limiting resource for striped mice (likely due to being 117 available year-round in succulent shrubs, Schoepf *et al.* (2017), and precipitation 118 thus affects them primarily through the availability of annual food plants. 119

Density regulation is a key component in population dynamics of small mam-120 mals (Krebs, 2013), and we calculated a proxy for monthly population density by 121 dividing the number of trapped mice by the study area size. We only included 122 adult female mice in the measure, as immature individuals do not compete sig-123 nificantly for reproduction or food. Males were excluded to allow implementing 124 density feedback in the population model without having to make assumptions 125 about sex ratio in family groups and number of male floaters. Further, competi-126 tion within sexes can be much more important than between sexes (Wauters et al., 127 2004). This modelling decision was unproblematic as vital rate model selection and 128 matrix model predictions were not sensitive to inclusion of the males in the density 129 measure (results not shown). As a measure for population density was required as 130 a covariate for the mark-recapture model, it was not possible to correct the counts 131

of trapped mice with recapture probabilities at that stage. However, due to overall high and invariable recapture probabilities over the study period, the raw counts are representative of the total population size (Nater *et al.*, 2016a). The size of the study site varied throughout the study period due to changes in the size of individual home ranges (Schradin *et al.*, 2010), and we estimated it using a 100% minimum convex polygon (MCP) approach (Worton, 1987) on the coordinates of the sampled nests in each month.

#### **139 ADAPTATION OF VITAL RATE ESTIMATION**

In a previous study (Nater et al., 2016a) we had estimated several monthly vital 140 rates of female striped mice as functions of ambient temperature, food availability, 141 and population density for the period from January 2005 to September 2014. 142 Specifically, we had used (1) a multi-state mark recapture model (Lebreton *et al.*, 143 2009) to estimate monthly survival and maturation probabilities of immatures 144  $(S_i, \Psi_{ib})$  and philopatrics  $(S_p, \Psi_{pb})$ , and survival probability of breeders  $(S_b)$ , (2) a 145 generalized linear mixed model (GLMM) to estimate the breeding probability (B)146 of breeders and (3) GLMs to estimate litter probability (L) and litter size (F). For 147 using the vital rate - environment relationships in a matrix population model in 148 this study, we re-defined the breeder stage to accommodate a pure post-breeding 149 census and elevated litter size using auxiliary data to obtain population projections 150 that did not go extinct within a few months. These adaptations and the resulting 151 re-analyses of vital rate - environment relationships are detailed in Appendix 1. 152

#### 153 STAGE-STRUCTURED POPULATION MODEL

We used the re-estimated vital rate - environment relationships to build a stagestructured population model for the female segment of the striped mouse population. We defined population structure at time t (N(t)) as a vector containing the number of immatures ( $N_i(t)$ ), philopatrics ( $N_p(t)$ ) and breeders ( $N_b(t)$ ):

$$N(t) = \begin{bmatrix} N_i(t) \\ N_p(t) \\ N_b(t) \end{bmatrix}$$

We then defined transition matrices A(t) (for more information see Caswell, 2001) that describe the monthly transitions between these stages depending on the vital rates in striped mouse life cycle (Figure 1). As all vital rates were functions of monthly temperature, food availability, and population density, the matrix itself was dependent on these environmental covariates (indicated by time-dependence of A):

$$A(t) = \begin{bmatrix} S_i \Psi_{ib} LF & S_p \Psi_{pb} LF & S_b BLF \\ S_i (1 - \Psi_{ib}) & S_p (1 - \Psi_{pb}) & 0 \\ S_i \Psi_{ib} & S_p \Psi_{pb} & S_b \end{bmatrix}$$

154

This allowed projecting population size and structure from a given month t to the next (t+1) using N(t+1) = A(t)N(t).

157

<sup>158</sup> We assessed the ability of the matrix model to capture striped mouse popula-<sup>159</sup> tion dynamics by comparing model-generated population hindcasts to the observed

population sizes over the course of the study period. We used the observed num-160 bers of immatures, philopatrics and breeders at the beginning of the time series to 161 define the initial population vectors N(t = 1) and N(t = 2). We then projected the 162 population for 116 time steps using projection matrices A(t) generated with the 163 observed time-series of temperature and food availability. The density covariate 164 was calculated from the projected population size N(t) at every timestep (details 165 in Appendix 2), thus letting population density propagate within the model. We 166 ran one projection using the specific year random effect values estimated by the 167 breeding probability model and another 100 trajectories by sampling this random 168 effect from a normal distribution with the estimated variance. Subsequently, we 169 compared the monthly population numbers predicted by matrix model projections 170 to the observed number of trapped mice in each month. 171

#### 172 RANDOM DESIGN LTRE

Population dynamics show responses of varying magnitude to changes in differ-173 ent vital rates and the environmental drivers underlying these changes (Oli, 2004; 174 Coulson et al., 2000). Life Table Response Experiments (LTREs) are retrospective 175 perturbation tools for quantifying relative impacts of matrix elements, vital rates 176 and lower-level covariates on previously observed population dynamics (Caswell, 177 2001). The dominant right eigenvalues of the matrices ( $\lambda$ ) is often used as the re-178 sponse variable of interest in LTREs as it represents long-term population growth 179 rate for density-independent populations in constant environments (Caswell, 2001). 180 Despite this definition not applying here,  $\lambda$  was highly correlated with transient, 181 one-time-step growth rate in our model (Appendix 4: Figures S4.1 & S4.2), and 182 we thus used it as the response variable of an LTRE analysis (for results using 183

transient one-time-step growth rate instead, see Appendix 3). As we had 116 dif-184 ferent matrices available (one for each month of the study period), we used a 185 random design LTRE (Caswell, 2001, chapter 10.2) to decompose temporal vari-186 ation in  $\lambda$  into contributions from variation in all different vital rates and from 187 the changes in temperature, food availability and population density underlying 188 it. This required sensitivity estimates of  $\lambda$  to changes in environmental covariates, 189 vital rates and matrix elements, and we calculated these numerically using the 190 element-by-element mean of all 116 matrices as a reference (Horvitz et al., 1997). 191 We performed the random design LTRE analysis for all matrices together, as well 192 as pooled into three seasons: breeding season (Aug - Nov), dry season (Dec - Mar) 193 and cold season (Apr - Jul). 194

#### 195 QUALITATIVE HINDCAST PERTURBATION

Projections from our matrix model were characterized by population peaks of 196 varying frequency and amplitude (Figure 2). To gain insight into the roles of envi-197 ronmental covariates in producing these patterns in our model, we did a qualitative 198 hindcast perturbation analysis. Specifically, we created 6 hindcast projections in 199 which we disabled the effects of one or two of the environmental drivers by set-200 ting them to a constant value, and compared these projections to the unperturbed 201 hindcast. The constant value chosen for all covariates was their mean over the 202 study period, and the random year effect on breeding probability was set to 0 203 here. 204

#### 205 STOCHASTIC DYNAMICS IN A STATIONARY ENVIRONMENT

Stochasticity in environmental conditions is ubiquitous and can strongly influence 206 the dynamics of populations (Tuljapurkar, 2013), particularly in combination with 207 with density-dependence (Boyce et al., 2006; Sæther, 1997). Understanding pop-208 ulation dynamics of species like striped mice that inhabit a variable environment 209 and are strongly density-regulated thus requires consideration of environmental 210 stochasticity. We analyzed population dynamics in a stochastic but stationary en-211 vironment with a prospective perturbation analysis. We generated 10,000 stochas-212 tic 200-year time series of temperature and food availability by randomly sampling 213 sequences of the covariates from all years we had data for (2005 - 2014). We defined 214 the year as starting in April, as the population is generally between maximum and 215 minimum densities in this month and each year then contains a distinctive popula-216 tion peak and trough. For each replicate time-series, we then created two perturbed 217 scenarios in which either temperature  $(S_T)$  or food availability  $(S_F)$  was increased 218 by 1%. We used unperturbed and perturbed environment time-series to simulate 219 future population trajectories, starting the projections using observed population 220 numbers and stage distributions for April 2005. For the stochastic projections, we 221 introduced a maximum litter size threshold of 5.8 (maximum value in the data) 222 to prevent unrealistic behavior of the exponential litter size model. 223

For each population projection we obtained this way, we determined population size and structure during the population maximum and minimum in each year. Subsequently we calculated the fold-changes of population size and structure, as well as quasi-extinction probability (threshold = 5 females), in scenarios  $S_T$  and  $S_F$  relative to the unperturbed scenario. Finally, we used additional simulations in a pathway analysis to determine which vital rates were primarily responsible for the calculated changes in annual minimum/maximum population sizes under scenarios  $S_T$  and  $S_F$ . We describe this analysis in more detail in Appendix 2.

#### 232 STOCHASTIC DYNAMICS IN A CHANGING ENVIRONMENT

When studying population responses to climate change, a gradual instead of a 233 constant change in temperature (and potentially other environmental factors) has 234 to be considered (Wolkovich et al., 2014). We explored how stochastic population 235 dynamics of striped mice changed when the population experienced a gradual in-236 crease in temperature. The magnitude of this temperature change was inspired 237 by climate change scenarios for South Africa. General Circulation Models predict 238 climate warming in the Succulent Karoo to range from 0.01°C to 0.4°C per year 239 until the end of the century, with the majority of models predicting and increase 240 of 0.02°C to 0.03°C per year (Jury, 2013). We thus assumed an annual tempera-241 ture increase of 0.025°C for our first climate change scenario  $(S_{T\uparrow})$ . It is unlikely 242 that the availability of food plants for striped mice will remain unchanged in the 243 future, as rainfall is generally expected to decrease. Climate change scenarios for 244 rainfall in South Africa are very variable and subject to large local differences 245 (MacKellar et al., 2007; Collier et al., 2008), however. This, in combination with 246 direct detrimental effects of increasing temperatures on Karoo vegetation (Musil 247 et al., 2009), makes quantitative predictions of vegetation changes very difficult. 248 We therefore adopted an exploratory approach regarding changes in food availabil-249 ity and included two additional climate change scenarios in which food availability 250 was expected to gradually decrease up to 10%  $(S_{T\uparrow F\downarrow})$  or 20%  $(S_{T\uparrow F\downarrow\downarrow})$  within 100 251 years. 252

For each climate change scenario, we generated 20,000 100-year stochastic en-253 vironment time series and perturbed them by imposing trends in temperature 254 and food availability as described above. We then ran population projections for 255 1200 time-steps using the original and perturbed environment time series, and an-256 alyzed changes in annual minimum/maximum population sizes and population 257 structures. Contrary to the analyses on stationary environments, we here not 258 only looked at the correlation between the original and perturbed trajectories, 259 but also accounted for temporal changes by fitting a linear model of the form 260  $X_{perturbed} \sim X_{original} + year + year : X_{original}$  where X represents the quantity of 261 interest (population size or structure). For each simulation in each scenario, we 262 used the parameters of this linear model to make an estimate of the fold-change 263 in X after 100 years of exposure to the climate change scenario. The "original" 264 population size and structure we used in those calculations were based on the 265 mean value for the entire unperturbed scenario for each simulation. In a last step, 266 we quantified the contributions of different demographic pathways to changes in 267 annual minimum/maximum population sizes after 100 year exposure to climate 268 change scenarios (details in Appendix 2). 269

<sup>270</sup> All matrix model analyses were done in R version 3.4.0 (R Core Team, 2015).

## 271 Results

#### 272 MODEL FIT & POPULATION GROWTH RATES

A model hindcast using the observed levels for the random effects in breeding probability performed well at capturing observed population dynamics (Figure 2). The timing of population increases and decreases was well represented by the model, <sup>276</sup> but peak population sizes tended to be somewhat overestimated. Furthermore, we
<sup>277</sup> demonstrated that even when the values of the random year effects on breeding
<sup>278</sup> probabilities were sampled randomly, the predicted pattern was well preserved.

We calculated two measures of population growth rate for each of the 116 ma-279 trices. The dominant right eigenvalues of the time-specific matrices ranged from 280 0.82 to 1.6 (mean = 1.01, SD = 0.18). Transient one-time-step growth rates (calcu-281 lated using both time-specific matrices and observed population structures) took 282 values between 0.81 and 1.95 (mean = 1.06, SD = 0.25). Dominant right eigenval-283 ues and transient growth rate followed the same pattern and were numerically very 284 similar (Appendix 4: Figure S4.1). 93% of variation in transient growth rates was 285 explained by the dominant right eigenvalues of the matrices (Appendix 4: Figure 286 S4.2). 287

# 288 RELATIVE STRENGTHS OF DRIVERS OF PAST POPULATION289 CHANGE

Decomposition of variation in the dominant right eigenvalues of the matrices using 290 a random design LTRE identified food availability, working primarily through lit-291 ter size, as the main driver of past population changes. The contribution of changes 292 in food availability was 0.021, and thus three times as large as the next influential 293 quantity, variation in population density (contribution = 0.007). Contributions 294 from variation in temperature and covariation among different environmental fac-295 tors were comparatively small (Figure 3). Among vital rates, the strongest contri-296 bution came from variation in litter size (0.008), followed by variation in philopatric 297 maturation probability (0.005, Appendix 4: Figure S4.4). Other considerable con-298 tributions came from variation in breeding probability and litter probability, as 299

well as their covariances with each other and with philopatric maturation probability. These relative rankings were conserved when variation in transient growth rate (instead of dominant right eigenvalue) was the quantity of interest (Appendix 303 3).

Seasonal analysis revealed that food availability only had the largest contributions in the breeding and the cold season. In the dry season, the largest contribution was ascribed to population density (Appendix 4, Figure S4.3). Similarly, contributions from philopatric maturation probability were more important than contributions from litter size in the dry season only (Appendix 4, Figure S4.5).

#### **309 QUALITATIVE PERTURBATION OF POPULATION HINDCASTS**

Excluding temporal variation in environmental covariates led to different changes 310 in population dynamics (Figure 4). Removing temperature variation resulted in 311 only slight changes in the height of population peaks. When the effect of food 312 availability was ignored, on the other hand, both population increases and de-313 creases were lost almost completely, irrespective of whether or not temperature 314 was allowed to vary. Finally, disabling the density feedback led to rapid population 315 explosion. Again, this was the case both with varying and constant temperature. 316 Disabling the density feedback in the presence of constant food availability re-317 sulted in a population that was almost constant at a small size, and showed only 318 low amplitude fluctuations as a consequence of varying temperature. 319

#### 320 STOCHASTIC DYNAMICS

<sup>321</sup> Population trajectories obtained for stochastic environments were characterized
<sup>322</sup> by marked differences in peak population sizes (Appendix 4: Figure S4.6), but

every peak was followed by a similar crash. This behavior led to perturbed trajectories that differed from original trajectories in minimum and maximum annual population sizes, but did not diverge strongly from original trajectories over time (Appendix 4: Figure S4.7).

Population peaks in stationary environments became higher when either tem-327 perature (mean fold-change = 1.009) or food availability (mean fold-change =328 1.025) was increased by 1% (Figure 5a & b). In both perturbations, these increases 329 were primarily driven by changes in litter size and the maturation probability of 330 immatures (Figure 5c & d). Minimum population sizes became larger under in-331 creased food availability (mean corr. coefficient = 1.005) due to changes in litter 332 probability, breeding probability and philopatric maturation probability. Increased 333 temperature, on the other hand, could lead to either smaller or larger minimum 334 population sizes depending on the stochastic sequence of years, and contributions 335 were spread over several vital rates (Figure 5b & d). Perturbations also affected 336 population structure: during population peaks and lows, higher temperature led to 337 a larger proportion of philopatrics, while higher food availability resulted in more 338 philopatrics and breeders (Appendix 4: Table S4.1). Extinction probability over 339 the simulated 200-year period was small at 5.2% for unperturbed trajectories, but 340 decreased further to 4.8% and 4.9% when temperature and food availability were 341 increased. 342

Annual maximum and minimum population sizes displayed distinct responses to different climate change scenarios (Figure 6a & b). With gradually increasing temperature but no change in food availability  $(S_{T\uparrow})$ , peak population sizes increased by 11.2%, while minimum population sizes decreased by 2.6% on average over a 100-year period. This positive effect of increasing temperature on

peak population sizes was offset if accompanied by a 10 % gradual decrease in 348 food availability  $(S_{T\uparrow F\downarrow})$ : Maximum population size after 100 years was lower in 349 all simulations, with an average decrease of 10.1%. Minimum population size de-350 creased by an average of 4.2%, although increases were seen in some simulations 351 (Figure 6b). When food availability decreased even more (20% in  $S_{T\uparrow F\downarrow\downarrow}$ ), maxi-352 mum population size decreased by 29.6% and minimum population size by 12.7%353 on average. Decreases in minimum population sizes were driven by changes in mat-354 uration, breeding and litter probabilities. The same vital rates were responsible for 355 lower peak population sizes under  $S_{T\uparrow F\downarrow}$  and  $S_{T\uparrow F\downarrow\downarrow}$ , while the higher population 356 peaks under  $S_{T\uparrow}$ ) were primarily due to changes in litter size. 357

Stage structure during the maximum and minimum population sizes was affected similarly by all three climate changes scenarios: Populations had a considerably higher proportion of philopatrics and lower proportion of breeders after 100 years of exposure to the scenarios (Appendix 4: Figure S4.8 & Table S4.2). The proportion of immatures on the other hand decreased only when food availability was projected to go down  $(S_{T\uparrow F\downarrow} \& S_{T\uparrow F\downarrow\downarrow})$  and showed very variable responses when only temperature increased  $(S_{T\uparrow})$ .

Extinction probability was almost halved when temperature increased gradually (decrease from 2.6% without perturbation to 1.4% under  $S_{T\uparrow}$ , fold decrease = 1.83). An accompanying moderate decrease in food availability ( $S_{T\uparrow F\downarrow}$ ) resulted in a 1.29-fold decrease in extinction probability (2.6 to 2.0%), whereas with a higher food decrease ( $S_{T\uparrow F\downarrow\downarrow}$ ), it increased 1.45-fold (2.6 to 3.8%).

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## 371 Discussion

In this study, we analysed population dynamics of a small semi-desert rodent on multiple temporal scales ranging from short-term month-by-month changes to long-term projections spanning several decades. By explicitly including environmentdemography relationships, we obtained a population model able to reproduce past population dynamics and identified food availability affecting reproduction and resulting density feedbacks as the main mechanisms driving the strongly fluctuating population dynamics of our study species.

Analysing the population dynamics on short timescales, we found that variation 379 in monthly population growth rates was mainly due to changes in food availability 380 (Figure 3) affecting vital rates linked to reproduction (predominantly litter size, 381 Appendix 4: Figure S4.4). The same general pattern emerged for multi-annual 382 stochastic population dynamics: elevating food availability resulted in larger pop-383 ulation sizes during annual peak and low phases as a result of changes in matura-384 tion rates and reproductive output (Figure 5). The importance of food availability 385 was emphasized further by the fact that population fluctuations largely disap-386 peared when variation in plant cover was ignored (Figure 4). These findings are 387 consistent with other studies that found food availability to be the main driver 388 of population fluctuations of rodents in semi-arid environments (Brown & Ernest, 389 2002; Previtali et al., 2010; Lima et al., 2008), as well as primary consumers in 390 general (Hunter & Price, 1992; Kagata & Ohgushi, 2006). In the case of striped 391 mice, opportunistic breeders displaying a high degree of plasticity in reproductive 392 timing (Raynaud & Schradin, 2008; Nel et al., 2015), the population increases fol-393 lowing elevated food availability are likely the results of prolonged reproductive 394

seasons with many young animals starting to breed early, potentially followed by
"out-of-season" reproduction due to more favorable conditions.

The second-most important driver of changes in monthly population growth 397 rates was population density (Figure 3). Variation in population growth rates dur-398 ing months with scarce food (dry season) was even primarily due to changes in 399 population density (Appendix 4: Figure S4.3). Stronger density feedbacks when 400 populations are close to their carrying capacity (e.q. due to resource scarcity) have 401 been found in a range of species including other rodents (Goswami et al., 2011), un-402 gulates (Albon et al., 2000; Coulson et al., 2001), and raptors (Krüger, 2007). This 403 reflects the importance of population density in regulating and stabilizing popu-404 lation dynamics (Hanski, 1990), and is further supported by our model predicting 405 either rapid population explosion or extinction when the density feedback was 406 disabled (Figure 4). The impacts of density feedbacks were very evident in multi-407 annual population dynamics as well: While both increases in food availability or 408 temperature led to larger annual minimum and maximum population sizes (Figure 409 5a & b), these changes did not add up over time and populations in perturbed en-410 vironments did not progressively diverge from those in unperturbed environments 411 (Appendix 4: Figure 4.7). This indicates that density regulation is strong enough to 412 "reset" population dynamics every year, and supports the hypothesis that strong 413 density feedbacks buffer populations of fast-living species against environmental 414 change (Williams, 2013). 415

Temperature only explained a small fraction ( $\approx 4\%$ ) of the variation in shortterm population growth rates (Figures 3) and increasing it led to smaller changes in long-term population abundances than increasing food availability (Figures 5). This is a result of relatively weaker impacts of temperature on vital rates (Nater *et al.*, 2016a), and may be partially related to the generally lower interannual
variation in temperature relative to food availability and population density.

Long-term population responses of striped mice under climate change scenarios 422 depended strongly on the assumed change in food availability. Rising temperatures 423 alone resulted in larger annual population peaks as a consequence of increased lit-424 ter size (particularly outside the main breeding season), and variable responses of 425 annual minimum population size (Figure 6). Adding a gradual decrease in food 426 availability, however, led to reduced maturation, breeding and litter probabilities, 427 and ultimately populations with overall lower abundance (minimum and maxi-428 mum population sizes, Figure 6). The strength of this response scaled with the 429 magnitude of the food decline. This lower reproductive output during the main 430 breeding season led to lower peak densities, which in turn let more philopatrics sur-431 vive the dry season and thus contribute to the following minimum population size. 432 This is supported by the generally higher percentage of philopatrics in populations 433 experiencing climate change (Appendix 4: Figure S4.8). Striped mice are known 434 to delay reproduction, likely to increase their chances of surviving the harsh dry 435 season and then breed the following spring (Schradin et al., 2012). Our analyses 436 indicate that under harsher conditions (e.g. increasing temperatures and decreas-437 ing food availability associated with climate change), more mice may be forced 438 to adopt this strategy, and possibly for a longer period of time. Delayed maturity 439 under adverse environmental conditions has been studied in long-lived species (e.q. 440 seabirds, Nevoux et al. (2010)), but less in short-lived species, making the striped 441 mouse an interesting system for future study of the role of delayed reproduction 442 in mediating environmental change. 443

444 The

The quasi-extinction probability of striped mouse populations over 100 years

increased only under the scenario with the strong decrease in food availability 445  $(S_{T\uparrow F\downarrow\downarrow})$ , and even then was still quite low at 3.8%. This too is related to the strong 446 density feedback stabilizing the population and thus buffering it even against large 447 changes in the environment. Under the investigated scenarios, our simulations thus 448 do not indicate that gradual climate change threatens persistence of striped mice 449 in the Succulent Karoo. However, the climate change scenarios we used here may 450 be only partially representative of future environmental change, and future studies 451 could improve on this from two perspectives. 452

First, the simulated decreases in vegetation cover were only qualitatively linked 453 to predicted future changes in rainfall and scenarios may therefore not accurately 454 represent the expected change in food availability due to climate change. Hav-455 ing a model linking vital rates directly to food availability - instead of indirectly 456 through rainfall as done in many other studies on arid ecosystems (e.q. Ozgul et al., 457 2014) - granted us more insight into the biological mechanisms underlying popu-458 lation dynamics. However, the downside of this is that in order to run this model 459 under more realistic climate change scenarios, the relationship between rainfall 460 and vegetation cover needs to be quantified. This is a non-trivial task, as vegeta-461 tion responds to rainfall in a complex and non-linear way, depending not only on 462 amount but also timing of rainfall (e.g. 100 mm of rain over the course of a month)463 stimulating plant growth vs. 100mm of rain falling within one day and destroying 464 vegetation). Second, climate change will not only lead to gradual changes in mean 465 environmental variables, but also increases in the frequency of extreme weather 466 events (IPCC, 2014). We have not considered such events in the current study, 467 but they can potentially have strong impacts on population dynamics (van de Pol 468 et al., 2010). In semi-arid Chile, for example, extreme flooding events following 469

catastrophic rainfalls can reset long-term demographic trends and community dy-470 namics (Kelt, 2011), and similar flooding events can occur in the Succulent Karoo 471 with potentially detrimental consequences for striped mice. Extreme events also 472 happen in the form of droughts and temperature anomalies, such as in the un-473 usually dry winter of 2003 which wiped out almost our entire study population 474 (pers. observation). So while the results from this study indicate that moderate 475 trends in average temperature and food availability do not threaten viability of 476 striped mouse populations, the same may not be true for changes in frequency and 477 intensity of extreme weather events. Quantifying the general relationship between 478 vegetation cover and rainfall, as well as occurrence and consequences of extreme 479 events (e.g. floods, droughts), are thus important topics for future research and will 480 allow us to make more realistic predictions of striped mouse population responses 481 to climate change. 482

Another area to improve on is the accuracy of numerical predictions. While our 483 population model produced a good fit to observed data, achieving this numerical 484 accuracy required adjustment of litter size estimates using auxiliary information. 485 The original measure of litter size had been consistently too low due to a combina-486 tion of low detectability of immatures and potential overestimation of the number 487 of breeding females (Appendix 1, Section 1.2). In order to obtain more accurate 488 numerical predictions, future studies should aim to quantify litter size more pre-489 cisely. This could be done, for example, by using open population mark-recapture 490 designs to account for the low detectability of immatures (Pradel, 1996), larger-491 scale genetic studies (sensu Schradin et al. (2012)) to reliably identify the females 492 giving birth and their offspring, and experiments investigating the environmental 493 effects on both the actual number of pups born to a female as well as nest survival 494

<sup>495</sup> of those pups.

Lastly, and particularly when discussing responses to climate change, one has 496 to keep in mind that in this study, we have considered only a single population 497 of striped mice in isolation, ignoring immigration and spatial dynamics and ac-498 counting for predation only indirectly (through survival estimates, Nater et al. 400 (2016a)). However, interactions and movement between different striped mouse 500 populations are likely important, and so is the general role of striped mice in the 501 foodweb. In (semi-)arid environments, small mammals often have important roles 502 as keystone (prey) species and ecological engineers (Kelt, 2011). If climate change 503 reduced overall abundance of striped mice (and potentially other rodent species) -504 as predicted in this study - this could have cascading effects on the abundance of 505 bird, reptile, and carnivore predators (Byrom et al., 2014), on the spread of inva-506 sive species (Madrigal et al., 2011), and on the plant species richness that makes 507 the Succulent Karoo a unique biodiversity hotspot (Hillebrand et al., 2007). Such 508 effects on other trophic levels could, in turn, feed back again on rodent (meta-509 )population dynamics. To make realistic predictions on how the Succulent Karoo 510 as a whole (and similar biomes) will respond to climate change, an ecosystem ap-511 proach coupling climate to plant, rodent and predator populations and including 512 feedbacks between the different trophic levels will therefore be invaluable. 513

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Environmental factors and density feedbacks have long been recognized as the key determinants of population dynamics, and studying these relationships via variation in vital rates has emerged as a powerful approach (Gamelon *et al.*, 2017). Nonetheless, environmental factors are rarely included into population models explicitly and few models consider interactions between the extrinsic environment

and density feedbacks (Ehrlén et al., 2016). In this study, we have accounted for 520 these complexities by building a density-dependent population model based on 521 estimated environment-vital rate relationships. Using this model, we have shown 522 that both short- and long-term population fluctuations of a semi-desert rodent are 523 sensitive to changes in food availability affecting reproduction, but also strongly 524 mediated by intrinsic feedbacks. Strong density dependence thus buffers this popu-525 lation against environmental change, and the environmentally explicit population 526 model enabled us to gain unique insights into the demographic mechanisms un-527 derlying this buffering. 528

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### <sup>536</sup> Author's contributions

<sup>537</sup> CN, KB, CC, CS and AO conceived the ideas and designed methodology. CS <sup>538</sup> collected the data. CN and KB analysed the data and led the writing of the <sup>539</sup> manuscript. All authors contributed critically to the drafts and gave final approval <sup>540</sup> for publication.

## 541 Data accessibility

<sup>542</sup> Data available from the Dryad Digital Repository: https://doi.org/10.5061/
<sup>543</sup> dryad.c1657.2 (Nater *et al.*, 2016b).

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## 753 Supporting information

- 754 The following Supporting Information is available for this article online: Appen-
- 755 dices 1-4.

## 756 Figures



**Figure 1:** Life cycle of the African striped mouse (post-breeding census). S indicates survival probabilities,  $\Psi$  maturation probabilities, B the breeding probability, L the litter probability, and and F the litter size. Subscripts for life stages: i =immature, p = philopatric, b = breeder).



**Figure 2:** Observed population sizes (black, dashed), model projections using observed levels of random variation (blue), and 100 model projections where random effects were sampled from a normal distribution (green).



Figure 3: Relative contributions of variances and covariances of temperature, food availability, and population density to variation in the dominant right eigenvalue  $\lambda$ .



Figure 4: Original and perturbed model hindcasts, where one or two environmental covariates are set to be constant at their mean value. Dashed lines have constant temperature, grey lines have constant food availability, and thick lines have constant density (disabled feedback).



Figure 5: Panels a) and b) show the distributions of proportional change in maximum and minimum population sizes over 200 years when either temperature (orange) or food availability (green) was increased by 1%. Panels c) and d) show the relative contributions of different demographic pathways to those changes. Colored bars represent mean values, black margins mark the 95% confidence interval. Estimates of both population size change and vital rate contributions are based on 10,000 replicates for each scenario.



Figure 6: Panels a) and b) show the distributions of proportional change in maximum and minimum population size after a 100-year projection under the three climate change scenarios. Panels c) and d) show the relative contributions of different demographic pathways to those changes. Colored bars represent mean values, black margins mark the 95% confidence interval. Estimates of both population size change and vital rate contributions are based on 20,000 replicates for each scenario.