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### Ontogenetic development underlies population response to mortality [post-print]

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1    Ontogenetic development underlies population response to mortality

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36 asymmetry, stage-structure.

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## ABSTRACT

Understanding demographic responses to mortality is crucial to predictive ecology. While classic ecological theory posits reductions in population biomass in response to extrinsic mortality, models containing realistic developmental change predict the potential for counterintuitive increase in stage-specific biomass, i.e., biomass overcompensation. Patterns of biomass overcompensation should be predictable based on differences in the relative energetic efficiencies of juvenile maturation and adult reproduction. Specifically, in populations where reproduction is the limiting process, adult-specific mortality should enhance total reproduction and thus juvenile biomass. We tested this prediction by inducing an array of stage-specific harvesting treatments across replicate populations of *Daphnia pulex*. In accordance with reproductive regulation, the greatest biomass response occurred in the juvenile *Daphnia* stage and this response occurred most strongly in response to adult mortality. Nevertheless, we failed to detect significant biomass overcompensation and instead report largely compensatory effects. In total, our work demonstrates that knowledge of population structure is necessary to accurately predict population dynamics, but cautions that further research is needed to illuminate the factors generating over-compensatory versus compensatory responses across natural populations.

## INTRODUCTION

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Understanding how populations respond to perturbations, such as changes in mortality or resource productivity, is central to predicting ecological dynamics (Nicholson 1957, Turchin 2003, Yang et al. 2008). Our understanding of such responses traditionally derives from unstructured models which assume reproduction and mortality are the only processes underlying population change. Yet, ontogenetic development is ubiquitous within natural populations (Werner and Gilliam 1984, Rudolf and Lafferty 2011, Miller and Rudolf 2011), and both reproductive output and mortality risk change as individuals grow (i.e., size- or stage-dependent life history traits; de Roos and Persson 2013). Accordingly, theory incorporating size or stage-dependence produces an array of population dynamic patterns not possible in analogous unstructured models, including biomass overcompensation, cohort cycles, emergent facilitation, and alternative stable community states (de Roos et al. 2003, Miller and Rudolf 2011, de Roos and Persson 2013, Huss et al. 2013, Schröder et al. 2014). While developmental variation has been incorporated into increasingly complex community models (van de Wolfshaar et al. 2006, Ohlberger et al. 2012, Leeuwen et al. 2014, Toscano et al. 2017), some of the most fundamental and far-reaching predictions of this new theory, such as population responses to perturbation, still require focused experimental testing. These tests are necessary to revise our understanding of the basic processes driving population and community dynamics.

Perhaps the most important and far-reaching way in which unstructured and structured model predictions diverge is regarding population responses to extrinsic mortality (de Roos et al. 2007), a widespread natural perturbation. Unstructured models generally predict reductions in population density and biomass due to mortality, and this basic prediction is implicit in the vast majority of classical ecological theory (but see (Abrams 2009, Schröder et al. 2014) for scenarios

76 in which unstructured models can produce positive responses to mortality, or “hydra effects”). In  
77 contrast, structured models containing a realistic representation of the developmental process  
78 predict, counterintuitively, that biomass can increase in response to mortality (de Roos and  
79 Persson 2013, Schröder et al. 2014). The ultimate cause of such positive biomass responses is  
80 ontogenetic asymmetry, or differences in the relative energetic efficiencies of developmental  
81 stages (de Roos et al. 2007). For example, for fish populations, competition is generally more  
82 intense within the reproductive adult stage versus within the juvenile stage (i.e., juveniles use  
83 energy more efficiently), a scenario termed reproduction regulation (Persson and de Roos 2006,  
84 Schröder et al. 2009). Here, biomass builds up within the adult stage due to a reproductive  
85 bottleneck at normal background mortality levels, and increasing adult mortality is expected to  
86 enhance juvenile fish biomass via increased reproduction. More generally, mortality acts to  
87 release the energetically less efficient (i.e., regulatory) developmental stage from intra-stage  
88 competition, thereby enhancing the biomass production of this stage. Thus, positive effects of  
89 mortality on stage-specific biomass should be predictable based on the ontogenetic asymmetry of  
90 a population (Persson and de Roos 2013). Empirical tests of such predictions can illuminate the  
91 importance of developmental variation in governing natural population dynamics.

92         The magnitude of positive biomass responses to mortality within structured populations  
93 should further depend on the stage-specificity of mortality. Using a general structured consumer-  
94 resource model, de Roos et al. (2007) show that stage-specific biomass overcompensation is a  
95 widespread response to mortality in populations characterized by ontogenetic asymmetry:  
96 overcompensation occurs in response to both stage-independent and stage-specific mortality,  
97 even when stage-specific mortality is induced within the non-regulatory stage (i.e., the stage  
98 exhibiting overcompensation). Nevertheless, the magnitude of overcompensation varies across

99 these different types of mortality: overcompensation is greatest when mortality targets the  
100 regulatory stage, intermediate when mortality is stage-independent, and least when mortality  
101 targets the non-regulatory stage (de Roos et al. 2007). Though empirical studies testing for  
102 biomass overcompensation are still rare (Schröder et al. 2014), only one study to our knowledge  
103 has induced (stage-specific) mortality across different stages while monitoring biomass, rather  
104 than density, responses (Schröder et al. 2009, 2014). Such studies are ultimately necessary to  
105 pinpoint ontogenetic asymmetry in energetics as the property that allows for biomass  
106 overcompensation (Schröder et al. 2014).

107         The goal of our study was to test for the occurrence of biomass overcompensation, a key  
108 population dynamic phenomenon, as driven by developmental change. To accomplish this, we  
109 harvested different developmental stages within lab populations of *Daphnia pulex* while  
110 monitoring stage-specific and total population biomass and density responses over 7 *Daphnia*  
111 generations. Evidence for ontogenetic asymmetry within *Daphnia* enables us to make clear  
112 predictions regarding the effects of developmental variation on population dynamics. Intra-stage  
113 competition appears stronger within adult versus juvenile *Daphnia* (i.e., reproductive regulation:  
114 present study, Nilsson et al. 2010) and given this adult-dominated population structure, we  
115 hypothesized that positive biomass responses to mortality would occur in the juvenile, rather  
116 than the adult *Daphnia* stage. Furthermore, we predicted overcompensation would be strongest  
117 in response to adult, rather than juvenile *Daphnia* mortality, because adult mortality most  
118 directly reduces competition within this regulatory stage (de Roos et al. 2007). While our results  
119 support the predicted stage-specificity of positive biomass responses, we detected largely  
120 compensatory rather than overcompensatory biomass responses. Thus, our study demonstrates  
121 the importance of ontogenetic development in population responses to perturbation but cautions

122 that more work is needed before biomass overcompensation is presumed a general phenomenon  
123 in nature.

## 124 METHODS

### 125 *Study system*

126 We tested population responses to stage-specific mortality using replicate lab populations  
127 of *Daphnia pulex*. *Daphnia* (Order: Cladocera) is a key herbivore and important prey for size-  
128 selective predators within freshwater food webs (Brooks and Dodson 1965, Sarnelle 1993,  
129 Boersma et al. 1996, Wagner et al. 2004). A number of studies have explored the role of  
130 developmental processes in generating *Daphnia*'s well-known cyclic population dynamics (e.g.  
131 McCauley et al. 1990b, 2008), establishing this genus as an important model regarding the  
132 population dynamic consequences of ontogenetic stage structure.

133 We collected *Daphnia* from a small fishless pond in Huntsville, Texas and maintained  
134 populations under high food conditions in the lab for several months prior to experiments.  
135 Reproductively mature, female *Daphnia* produce clonal female offspring when environmental  
136 conditions are favorable and ephippia (haploid eggs that are fertilized by males) when resources  
137 are scarce. *Daphnia* size at reproductive maturity depends on food availability and temperature,  
138 among other environmental variables (McCauley et al. 1990a, Gurney et al. 1990, Stibor 1992).  
139 Within the present study (i.e., our lab environment), *Daphnia* reached reproductive maturity (i.e.,  
140 began producing eggs) as small as 0.9 mm. Thus, we considered *Daphnia* below this size  
141 threshold as juveniles and *Daphnia* above this size threshold as adults, though our findings do  
142 not depend on the precise threshold value (Supplementary Fig. 1).

### 143 *Treatments*

144 *Daphnia* populations within our experimental setup are dominated by adult biomass in  
145 the absence of extrinsic mortality (Fig. 2A, control treatment; see also Nilsson et al. 2010). For  
146 reasons explained in the *Introduction*, mortality inflicted on adults (relative to mortality targeting  
147 juveniles or stage-independent mortality) should produce the greatest positive biomass response  
148 because it most directly reduces competition within the adult stage (de Roos et al. 2007).  
149 Accordingly, our experimental design focused on adult harvesting.

150 We applied 4 treatments to replicate *Daphnia* populations (n = 4 replicates per  
151 treatment): (1) a high level of adult-specific harvesting (instantaneous adult mortality rate: 0.07  
152 per day); (2) a low level of adult-specific harvesting (instantaneous adult mortality rate: 0.04 per  
153 day); (3) a high level of juvenile-specific harvesting (instantaneous juvenile mortality rate: 0.07  
154 per day); and (4) a no-harvest control. Overcompensation occurs as a hump-shaped response  
155 between biomass and mortality (de Roos et al. 2007, Nilsson et al. 2010), and thus we harvested  
156 adult *Daphnia* at two different levels to increase our chances of capturing this hump-shaped  
157 region. Furthermore, a previous study on *Daphnia* biomass overcompensation induced a single  
158 high level of stage-independent mortality (instantaneous total mortality rate: 0.2 per day), which  
159 the authors suggested could have overwhelmed any positive biomass response (Nilsson et al.  
160 2010). Again, due to the hypothesized reproduction regulation of *Daphnia* populations, we  
161 excluded a low-level juvenile-specific harvesting treatment from our design because we expected  
162 juvenile mortality to induce the weakest biomass response.

### 163 *Experimental setup*

164 We ran the experiment in a semi-chemostat system. Our system used peristaltic pumps to  
165 deliver algae continuously to microcosms housing replicate *Daphnia* populations. Each  
166 cylindrical microcosm (3 L volume) featured four 63  $\mu$  nylon mesh-covered outflow holes at the

167 water surface to retain *Daphnia* but allow suspended particulate matter to pass through. This  
168 flow-through setup creates semi-chemostat resource dynamics, an assumption of models in  
169 which the conditions for biomass overcompensation have been explored (de Roos et al. 2008,  
170 Nilsson et al. 2010, Huss and Nilsson 2011). Microcosms (n = 16) were arranged in 2 adjacent  
171 spatial blocks and treatments were assigned randomly within each spatial block (2 replicates per  
172 block). Microcosms within each spatial block shared the same peristaltic pump, algae stock tank  
173 and drainage system.

174 We used a 3:1 mixture of natural and simulated pond water as the experimental medium  
175 because our previous work has shown this mixture effective in maintaining *Daphnia* populations  
176 over multiple generations. Natural pond water was collected monthly from a small pond in Sam  
177 Houston National Forest, Texas. This pond water was then stored in a refrigerated room prior to  
178 preparation for use in the experiment. Pond water preparation followed a 2-step purification  
179 process including 10  $\mu$  vacuum filtration and 3 h autoclave sterilization. Artificial pond water  
180 was created following the ‘animal medium’ recipe in Wyngaard and Chinnappa (1982). All  
181 experiments were conducted in a temperature-controlled room (set at 22° C) under a 12:12 dark-  
182 light cycle.

### 183 *Algae culture and delivery to microcosms*

184 The flagellated green alga *Chlamydomonas reinhardtii* was cultured in the lab and used  
185 as the food resource for *Daphnia*. Algae (CC-1010 wild type mt+ [UTEX 90], *Chlamydomonas*  
186 Resource Center) was grown in flasks containing TAP (Tris-Acetate-Phosphate) growth medium  
187 under continuous light. Algae was harvested near peak density and centrifuged at 5000 rpm for 8  
188 minutes. After centrifugation, TAP medium was discarded and replaced with simulated pond  
189 water. We then measured algae cell density using a hemocytometer and added this concentrated

190 algae to stock tanks containing 16 L of the experimental medium. We adjusted the volume of  
191 algae added to stock tanks to maintain a constant resource level ( $157,480 \text{ cells ml}^{-1}$ ) entering  
192 microcosms throughout the experiment. Stock tanks containing magnetic stir bars were  
193 continuously mixed over stir plates to keep algae suspended. Peristaltic pumps drew diluted  
194 algae from stock tanks and delivered it (using 1.42 mm ID Tygon tubing) to experimental  
195 microcosms ( $0.945 \text{ mL min}^{-1}$  inflow rate,  $0.019 \text{ h}^{-1}$  dilution rate) housing *Daphnia* populations.  
196 Algae stock tanks that fed microcosms were cleaned and replenished with fresh algae and  
197 experimental medium every 48 h throughout the duration of the experiment.

#### 198 *Experimental harvesting*

199 *Daphnia* were introduced to microcosms (50 individuals per microcosm) on 11  
200 December 2017 and populations were allowed to increase and enter a regular pattern of cycling  
201 before the start of harvesting treatments. *Daphnia* populations were harvested once per week  
202 from 16 January 2018 until 27 March 2018 (11 total harvesting events). *Daphnia* generation time  
203 within our semi-chemostat system is roughly 2 weeks and thus our experiment allowed for  
204 approximately 7 *Daphnia* generations.

205 During each harvesting event we first stirred microcosm contents to homogenize  
206 contents. Using a 50 mL Hensen Stempel zooplankton sampling pipette, we then removed either  
207 50% of the microcosm volume (1500 mL, high level harvesting treatments) or 25% of the  
208 microcosm volume (750 mL, low level harvesting treatment), depending on the assigned  
209 treatment. For the control treatment we removed 750 mL of medium containing *Daphnia* and  
210 followed the process for removing adults (see explanation below) but returned all contents to  
211 control microcosms (i.e., a sham treatment).

212 To remove adult *Daphnia*, water from microcosms was filtered through 530  $\mu$  mesh that  
213 retained adults but allowed juveniles to pass through. This mesh was then rinsed thoroughly to  
214 capture adult *Daphnia*. The water containing any *Daphnia* that passed through the mesh was  
215 returned to microcosms. To remove juveniles, water was first filtered through 530  $\mu$  mesh. This  
216 mesh was rinsed to capture adult *Daphnia*, which were returned to microcosms. The water  
217 containing juvenile *Daphnia* that passed through the 530  $\mu$  mesh was next filtered through 153  $\mu$   
218 mesh that retained juvenile *Daphnia* but allowed algae and resting eggs to pass through (also  
219 returned to microcosms). This 153  $\mu$  mesh was then rinsed to capture juvenile *Daphnia*. All  
220 *Daphnia* removed from microcosms were processed as detailed below (*Sampling*) to estimate the  
221 amount and stage-specificity of *Daphnia* biomass removed by harvesting treatments.

#### 222 *Sampling*

223 We sampled microcosms twice per week throughout the duration of the experiment.  
224 Sampling events occurred on the first and third days after harvesting. For each sample,  
225 microcosm contents were stirred, and 200 mL was removed using a 50 mL Hensen Stempel  
226 zooplankton sampling pipette. We measured the total length of the first 30 *Daphnia* individuals  
227 (or fewer if there were less than 30 individuals) within each 200 mL sample and counted eggs,  
228 young in brood pouches, and resting eggs for reproductive females. Any additional *Daphnia* (>  
229 30) within each 200 mL sample were counted. These samples were then returned to microcosms  
230 (i.e., non-destructive sampling).

231 We applied length-weight regressions to derive *Daphnia* biomass densities. If the total  
232 number of individuals per sample was less than 30, we applied a length-weight relationship  
233 (Nilsson et al. 2010) to individual length measurements to estimate juvenile, adult and total  
234 biomass per 200 mL sample. If the total number of *Daphnia* per sample was greater than 30

235 individuals, we used the size distribution from the 30 measured individuals to estimate the  
236 biomass of unmeasured individuals and combined these biomasses to yield biomass estimates per  
237 sample. To do this, we first calculated the proportion of individuals within 0.1 mm size classes  
238 (ranging from 0.3-1.2 mm) out of the 30 measured individuals within each sample. We then  
239 multiplied these proportions times the total number of unmeasured individuals, rounding down to  
240 the nearest individual. Assuming that unmeasured individuals within each size class were of the  
241 mean length (e.g., 0.45 mm in the 0.4-0.5 mm size class), we used these lengths to calculate the  
242 biomass of unmeasured individuals. We applied this same basic scaling procedure to derive  
243 estimates of stage-specific population densities per sample. Individuals  $\geq 0.9$  mm were  
244 considered adults and individuals  $< 0.9$  mm were considered juveniles in all calculations.

#### 245 *Analysis*

246 Our experiment consisted of 11 weeks of treatments and 27 sampling events between 5  
247 January 2018 and 13 April 2018. Two of the sampling events were conducted before treatments  
248 began and four were conducted after treatments ceased. Analyses covered the 21 sampling events  
249 during treatments and 2 sampling events after treatments had concluded to capture the effects of  
250 the final harvesting event.

251 We tested stage-specific harvesting treatment effects on: (1) the amount of stage-specific  
252 biomass removed from populations; (2) stage-specific and total population biomass and density;  
253 and (3) reproductive responses. We examined the amount of stage-specific biomass removed  
254 from populations to test the efficacy of harvesting treatments. We further tracked population  
255 density responses, in addition to biomass responses, to provide deeper insight into population  
256 responses to mortality (Schröder et al. 2014). Treatment effects on reproductive responses were  
257 examined to explore potential mechanisms behind biomass and density responses to mortality.

258 We calculated two reproductive response variables: average clutch size (a *per capita* measure)  
259 and total reproductive output (a population-level measure). Average clutch size was calculated  
260 by dividing the total number of eggs + offspring in brood pouches per sample by the number of  
261 ovigerous individuals per sample. Total reproductive output was calculated by multiplying  
262 average clutch size  $\times$  the proportion of ovigerous adults per sample  $\times$  total adult density per  
263 sample.

264 Auto-correlation function (ACF) plots indicated significant temporal autocorrelation in  
265 our response variables (measured as time series), violating the assumption of independence. To  
266 address this, we followed recommendations by (Zuur et al. 2009) to model autocorrelation  
267 structure within generalized least squares (nlme package [Pinheiro et al. 2018] in R software).  
268 For each response variable (i.e., each generalized least squares model), we utilized the most  
269 flexible autocorrelation structure: auto-regressive moving average, ARMA(p, q). We fit this  
270 autocorrelation model at the microcosm level (form = time | microcosm) within generalized least  
271 squares, obviating the need to model microcosm as a random effect. We further fit ARMA  
272 models to individual time series independently of generalized least squares to confirm these  
273 models did a good job of capturing *Daphnia* cycling.

274 All generalized least squares models included harvesting treatment, time and a treatment  
275  $\times$  time interaction as fixed effects. Biomass response variables were log-transformed, and density  
276 response variables were square-root transformed prior to model fitting. Our approach to finding  
277 an adequate ARMA(p, q) structure was to fit models with 1-7 total parameters (all possible  
278 combinations of p and q: 35 different structures) while retaining all fixed effects and select the  
279 model with the lowest AIC. When the best-fitting models were similar in AIC values ( $<2 \Delta$   
280 AIC), we chose the model with the fewest number of combined p, q parameters to maximize

281 parsimony. We note that according to (Zuur et al. 2009 and references therein), finding an  
282 adequate autocorrelation structure is sufficient to account for non-independence of data, with  
283 relatively little to be gained from finding a “perfect” fit. Here, the “adequate” p and q structure  
284 was the final autocorrelation structure for each response variable that captured most of the  
285 variance explained by temporal autocorrelation.

286 Each model was subjected to this same procedure to determine and incorporate the best  
287 fit ARMA(p, q) autocorrelation structure. Once an appropriate ARMA(p, q) structure was  
288 identified, we tested the overall significance of fixed effects by dropping these terms from  
289 models and comparing nested models using likelihood ratio tests. We tested for differences  
290 among harvesting treatment levels using least-squares means contrasts (lsmeans package (Lenth  
291 2016) in R software).

## 292 RESULTS

### 293 *Harvesting treatment efficacy*

294 Stage-specific harvesting treatments were largely successful in removing the targeted  
295 *Daphnia* developmental stage throughout the duration of the experiment (Fig. 1A). Though adult  
296 harvesting treatments (low-adult, high-adult) did remove some juvenile biomass, this was  
297 significantly less than that removed by the high-juvenile harvesting treatment (contrast, pooled  
298 adult harvesting treatments vs. high-juvenile:  $p < 0.001$ ; Fig. 1B). As intended, the high-adult  
299 harvesting treatment removed approximately twice the adult biomass of the low-adult harvesting  
300 treatment (pairwise contrast:  $p < 0.001$ ), while adult biomass removal in the high-juvenile  
301 treatment was negligible (Fig 1C). These treatment effects on juvenile and adult biomass  
302 removal were consistent throughout the duration of the experiment (likelihood ratio tests of  
303 treatment  $\times$  time interactions:  $p > 0.05$ ; Fig. 1A).

304 *Reproduction regulation*

305           In support of reproduction-regulation, adult *Daphnia* comprised 75% of total population  
306 biomass in the absence of harvesting (i.e., the control) when averaged over the duration of the  
307 experiment (Fig. 2A). In contrast, juveniles dominated total density, comprising 63% of all  
308 *Daphnia* in the control on average (Fig. 2B).

309 *Juvenile responses*

310           As hypothesized, juveniles exhibited stronger positive (though statistically insignificant)  
311 biomass and density responses to mortality compared to adults (biomass: Fig. 2C vs. 2E; density:  
312 Fig. 2D vs. Fig. 2F). We detected compensation, i.e., the lack significant differences between  
313 harvesting treatments and the control, in both juvenile biomass (contrast, pooled harvesting  
314 treatments vs. control:  $p = 0.718$ ; Fig. 2C, Fig. 3A, Fig. 3B) and juvenile density (contrast,  
315 pooled harvesting treatments vs. control:  $p = 0.612$ ; Fig. 2D). While the high-adult harvesting  
316 treatment increased juvenile biomass by 12% and density by 13% relative to the no-mortality  
317 control (Fig. 2C, Fig. 2D), these differences were not statistically significant (biomass, pairwise  
318 contrast:  $p = 0.258$ ; density, pairwise contrast:  $p = 0.178$ ). The only significant differences  
319 between harvesting treatment levels occurred between high-adult and high-juvenile treatments  
320 (biomass, pairwise contrast:  $p = 0.009$ ; density, pairwise contrast:  $p = 0.096$ ), which produced  
321 the most positive and most negative juvenile responses to harvesting, respectively (Fig. 2C, Fig.  
322 2D). All treatment effects on juvenile biomass and density were consistent throughout the  
323 duration of the experiment (likelihood ratio tests of treatment  $\times$  time interactions:  $p > 0.05$ ; Fig.  
324 3A, Fig. 3B).

325 *Adult responses*

326 In contrast, adult *Daphnia* biomass was significantly reduced by harvesting treatments  
327 when compared to the control (contrast, pooled harvesting treatments vs. control:  $p = 0.005$ ; Fig.  
328 2E, Fig. 3C, Fig. 3D). Reductions in adult biomass occurred across all harvesting treatment  
329 levels, including the treatment targeting juveniles (pairwise contrasts:  $p < 0.05$ ). Similar  
330 reductions due to harvesting occurred with *Daphnia* density (Fig. 2F), though this effect was not  
331 significant (contrast, pooled harvesting treatments vs. control:  $p = 0.114$ , Fig. 2B). These  
332 treatment effects on adult *Daphnia* biomass and density were consistent throughout the duration  
333 of the experiment (likelihood ratio tests of treatment  $\times$  time interactions:  $p > 0.05$ , Fig. 3C, Fig.  
334 3D).

#### 335 *Total population responses*

336 Because total *Daphnia* biomass was dominated by adults (Fig. 2A), total biomass  
337 responses to harvesting mirrored that of adult biomass: total biomass was significantly reduced  
338 by harvesting treatments when compared to the control (contrast, pooled harvesting treatments  
339 vs. control:  $p = 0.022$ , Fig. 2G). Total density was dominated by juveniles (Fig. 2B), and  
340 accordingly, showed compensation in response to harvesting treatments (contrast, pooled  
341 harvesting treatments vs. control:  $p = 0.450$ , Fig. 2E). Treatment effects on total *Daphnia*  
342 biomass and density were consistent throughout the duration of the experiment (likelihood ratio  
343 tests of treatment  $\times$  time interactions:  $p > 0.05$ , Fig. 3E, Fig. 3F).

#### 344 *Reproductive responses*

345 *Daphnia* responded to harvesting by increasing mean clutch size (*per capita* reproductive  
346 output) relative to the control (contrast, pooled harvesting treatments vs. control:  $p = 0.025$ ), with  
347 the high-juvenile harvesting inducing the greatest increase in clutch size (contrast, high-juvenile  
348 treatment vs. control:  $p = 0.027$ ) (Fig. 4A). The proportion of reproductive adults responded

349 similarly to harvesting treatments, though the effect of harvesting was marginal (contrast, pooled  
350 harvesting treatments vs. control:  $p = 0.0581$ ; Fig. 4B). These reproductive responses, when  
351 multiplied times the number of adults, resulted in compensation in total reproductive output  
352 across treatments (contrast, pooled harvesting treatments vs. control:  $p = 0.790$ ; Fig. 4C). These  
353 treatment effects on *Daphnia* reproductive responses were consistent throughout the duration of  
354 the experiment (likelihood ratio tests of treatment  $\times$  time interactions:  $p > 0.05$ ).

## 355 DISCUSSION

356 While it is generally assumed that extrinsic mortality (e.g., predation or disease) should  
357 reduce population density and thus biomass, this prediction derives from models that overlook a  
358 fundamental feature of natural populations: developmental variation. New theory incorporating  
359 food-dependent development instead predicts biomass increase in response to mortality (i.e.,  
360 biomass overcompensation) (Schröder et al. 2014), and biomass overcompensation underlies  
361 much of the higher-order, community-level consequences of developmental variation (Roos et al.  
362 2008, Huss and Nilsson 2011, de Roos and Persson 2013, Huss et al. 2014). *Daphnia* biomass  
363 responses to mortality in our study were largely consistent with those predicted by  
364 reproductively-regulated population theory, yet the magnitude of these effects was lower than  
365 expected (de Roos et al. 2007, Schröder et al. 2009). Specifically, juvenile *Daphnia* exhibited  
366 compensatory, but not over-compensatory, biomass responses to adult mortality (see also  
367 Nilsson et al. 2010). Thus, our work demonstrates that ontogenetic stage-structure mediates  
368 population response to extrinsic mortality, but cautions that further research is necessary to  
369 illuminate the factors underlying the occurrence of biomass overcompensation across natural  
370 populations.

371 *Testing theory: The juvenile response*

372 Our study explored biomass overcompensation within reproductively regulated *Daphnia*  
373 populations, which allowed us to test clear hypotheses regarding the importance of food-  
374 dependent development in mediating population response to mortality. Theory predicts that when  
375 populations are regulated by reproduction, adult mortality should enhance total reproduction and  
376 thus juvenile biomass because it reduces competition among adults. In line with this prediction,  
377 we found that juvenile *Daphnia*, the non-regulatory stage, exhibited the greatest compensation in  
378 biomass in response to mortality, with the strongest response induced by mortality of regulatory  
379 adults. As a consequence of compensation in juvenile biomass, mortality elicited a shift in the  
380 stage structure of *Daphnia* populations: The percentage of total biomass made up by juveniles  
381 increased from 25% (no extrinsic mortality control) to 32% (high adult harvesting treatment),  
382 and the percentage of juvenile individuals increased from 63% (no extrinsic mortality control) to  
383 71% (high adult harvesting treatment). All of these juvenile biomass and density responses were  
384 at least partially driven by a positive effect of mortality treatments on *per capita* adult *Daphnia*  
385 clutch size, leading to compensation in total reproductive output across treatments. Thus, we see  
386 broad correspondence between compensation in adult reproductive output and compensation in  
387 juvenile biomass and density in our study, though our analysis failed to detect differences in  
388 reproductive responses among mortality treatment levels.

389 While experiments inducing different types stage-specific mortality and measuring  
390 biomass responses are still exceedingly rare, our findings are consistent with other data from  
391 reproductively-regulated populations (Schröder et al. 2009, 2014, Nilsson et al. 2010). More  
392 specifically, the patterns we recovered regarding different types of stage-specific mortality match  
393 the most complete biomass overcompensation test to date on reproductively regulated poeciliid  
394 fishes (Schröder et al. 2009). Yet our findings differ from previous work in one main way: these

395 studies detected stage-specific biomass overcompensation in juveniles (Schröder et al. 2009,  
396 2014), while we detected biomass compensation (see also Nilsson et al. 2010). Notably, we  
397 detected juvenile biomass compensation across all mortality treatments, even when mortality was  
398 induced in the juvenile stage. We note that while biomass overcompensation in our experiment  
399 was statistically insignificant, high-level adult *Daphnia* harvesting did increase juvenile biomass  
400 by 12% and juvenile density by 13% relative to the no-mortality control. Regardless, while our  
401 experiment recovered patterns consistent with reproductively regulated population theory, we  
402 conclude that our effects are somewhat weak.

403         Several explanations exist for the lack of significant biomass overcompensation. First,  
404 *Daphnia* populations undergo multiple types of stage-driven population cycles, the physiological  
405 mechanisms behind which remain debated (McCauley et al. 2008, Martin et al. 2013, van der  
406 Meer 2016). Intrinsic *Daphnia* population cycles were clearly observed in our study, and most  
407 apparent in times series of the adult stage. While our analysis approach allowed us to account for  
408 intrinsic *Daphnia* cycling at the microcosm-level, such strong and complex temporal  
409 autocorrelation within *Daphnia* population dynamics could still obscure treatment effects.  
410 Second, while adult *Daphnia* in our study were reproductively mature, they could also continue  
411 to grow (see also Schröder et al. 2009). Thus adult *Daphnia* energy could be allocated to growth  
412 rather than reproduction, reducing the magnitude of any positive reproductive response  
413 (Ohlberger et al. 2011). However, we compared detailed *Daphnia* size distributions across  
414 treatment levels and found no evidence for more subtle shifts in size structure (e.g., within  
415 juvenile and adult stages; Supplementary Fig. 2). Third, it is possible that *Daphnia* do not exhibit  
416 overcompensation, or do so only under limited conditions. While earlier studies suggested  
417 biomass overcompensation in cladocerans (Slobodkin and Richman 1956, Edley and Law 1988),

418 a more recent study (Nilsson et al. 2010) instead showed juvenile biomass compensation in  
419 response to stage-independent (i.e., random) mortality. Lack of overcompensation in this study  
420 (Nilsson et al. 2010) could have occurred due to: (1) stage-independent mortality, which is  
421 expected to elicit weaker effects than stage-specific mortality (de Roos et al. 2007); or (2) the use  
422 of a single high mortality level that overwhelmed a potential overcompensatory response. We  
423 designed our experiment to include both low and high levels of stage-specific adult mortality in  
424 hopes of capturing a positive biomass response, and yet failed to do so. We did however find that  
425 juvenile biomass was 16% greater at the high versus low adult mortality level (though this effect  
426 was statistically insignificant), suggesting that an even higher mortality level could have  
427 captured a potential biomass response peak.

428         Still, both Nilsson et al. (2010) and the present study failed to demonstrate biomass  
429 overcompensation in *Daphnia*, raising the question of whether overcompensatory responses only  
430 exist within a limited parameter space (e.g., range of mortality levels) for certain taxa. Because  
431 overcompensation induces cascading effects of developmental variation on whole communities  
432 (de Roos and Persson 2013), weak effects might limit the importance of these responses for  
433 natural population dynamics and species interaction webs. Indeed, biomass overcompensation is  
434 the necessary prerequisite for several population and community effects of ontogenetic  
435 development (Roos et al. 2008, Huss and Nilsson 2011, de Roos and Persson 2013, Huss et al.  
436 2014), including emergent facilitation between predators (Roos et al. 2008, Huss et al. 2014) and  
437 alternative stable community states (van de Wolfshaar et al. 2006, Gårdmark et al. 2015,  
438 Toscano et al. 2016). Clearly, more empirical studies testing for biomass overcompensation in  
439 response to stage-specific mortality are needed before biomass overcompensation is presumed a  
440 general phenomenon in nature. Understanding the empirical factors that limit the magnitude of

441 biomass overcompensation in natural systems and incorporating these factors into theory is a  
442 major research priority moving forward.

443 *Testing theory: adult response*

444           Consistent with other experiments on reproductively-regulated populations (Schröder et  
445 al. 2009, Nilsson et al. 2010), adult biomass declined with mortality in our study. Thus, biomass  
446 compensation in our study was a stage-specific phenomenon exhibited only by juveniles. Still,  
447 we detected an interesting lack of treatment effects on adult biomass. From theory, we expected  
448 that: (1) high adult harvesting would reduce adult biomass the most; (2) high juvenile harvesting  
449 would reduce adult biomass the least, and; (3) low adult harvesting would have an intermediate  
450 effect on adult biomass. While our low adult harvesting treatment was successful in removing  
451 exactly half the adult biomass of the high adult harvesting treatment, we found no difference in  
452 long-term adult biomass between these treatments. Furthermore, harvesting juveniles produced  
453 the same reduction in adult *Daphnia* biomass, despite the juvenile mortality treatment being  
454 extremely accurate in capturing juveniles and not adults. One potential contributing factor to  
455 these results is that our adult harvesting treatments also removed some juvenile biomass.  
456 Regardless, these findings show that very different types of stage-specific mortality can produce  
457 a similar reduction in adult biomass. Thus, while juvenile biomass responses in our study were  
458 largely in line with reproductive regulation (de Roos et al. 2007, Schröder et al. 2009), the  
459 mechanisms underlying observed adult biomass responses require further study.

460 *Relevance for natural dynamics*

461           Our findings suggest that *Daphnia*, a key primary consumer within freshwater systems,  
462 exhibit remarkable resistance to extrinsic mortality. Despite inducing substantial stage-specific  
463 mortality (instantaneous mortality rate in high harvesting level treatments: 0.07 per day, or 50%

464 of stage-specific biomass removed per week), we found compensation in total *Daphnia*  
465 population density, suggesting that the increase in juveniles was roughly equal to the loss of  
466 adults through harvesting. While total *Daphnia* biomass was reduced by mortality due to the loss  
467 of large adults, juvenile biomass compensation suggests that size-selective predators feeding on  
468 adult *Daphnia* can help support predators feeding on juveniles (Brooks and Dodson 1965, Huss  
469 and Nilsson 2011). Testing this prediction under natural conditions could provide new insight  
470 into the role of zooplankton in supporting diverse predator guilds, and more broadly how  
471 ontogenetic development mediates energy flow through food webs (Reichstein et al. 2015).

## 472 *Conclusions*

473 Unstructured theory assuming individual equivalence has long dominated our  
474 understanding of population dynamics, yet ontogenetic variation is widespread and structured  
475 theory predicts fundamentally different responses to perturbation (de Roos and Persson 2013,  
476 Persson and de Roos 2013). At the most basic level, positive biomass responses arise due to the  
477 size-scaling of biomass production (e.g., via rates of food consumption, metabolism or mortality)  
478 (de Roos et al. 2007 Peters 1983, Werner and Gilliam 1984). Thus while our study used  
479 mortality to shift the adult-juvenile competitive balance, our findings extend to any type of  
480 perturbation that modifies ontogenetic asymmetry within populations (Schröder et al. 2014).  
481 Because such responses have clear applied relevance to the management of exploited populations  
482 (de Roos and Persson 2002, Gårdmark et al. 2015), as well as major implications for species  
483 interactions and community dynamics (Roos et al. 2008, Huss and Nilsson 2011, de Roos and  
484 Persson 2013, Huss et al. 2014), future experiments inducing a range of stage-specific mortality  
485 and measuring biomass responses are crucial. Such work could help to determine whether

486 biomass overcompensation as a result of food-dependent development should be incorporated as  
487 a core (i.e., default) feature of population and community theory moving forward.

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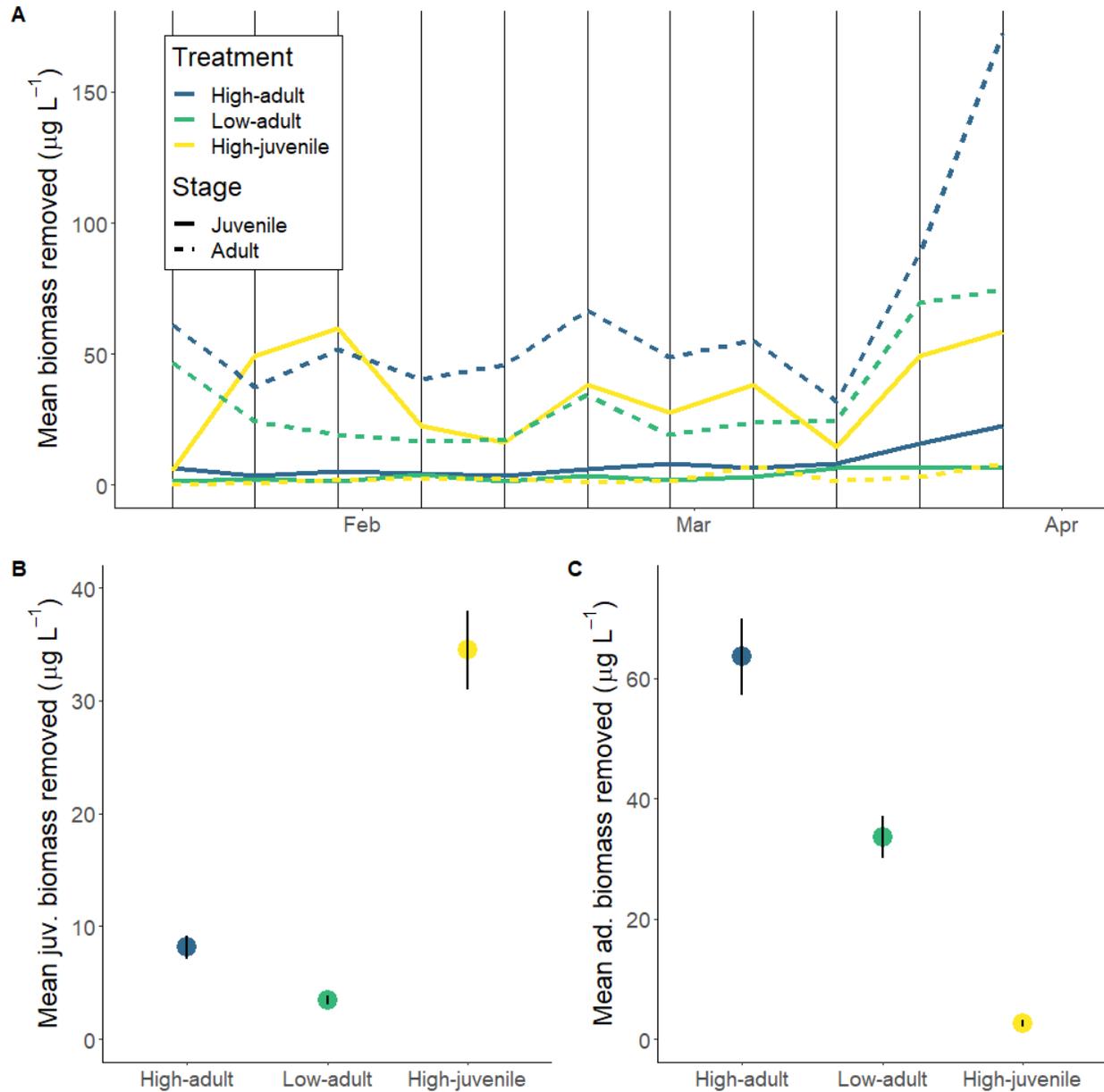
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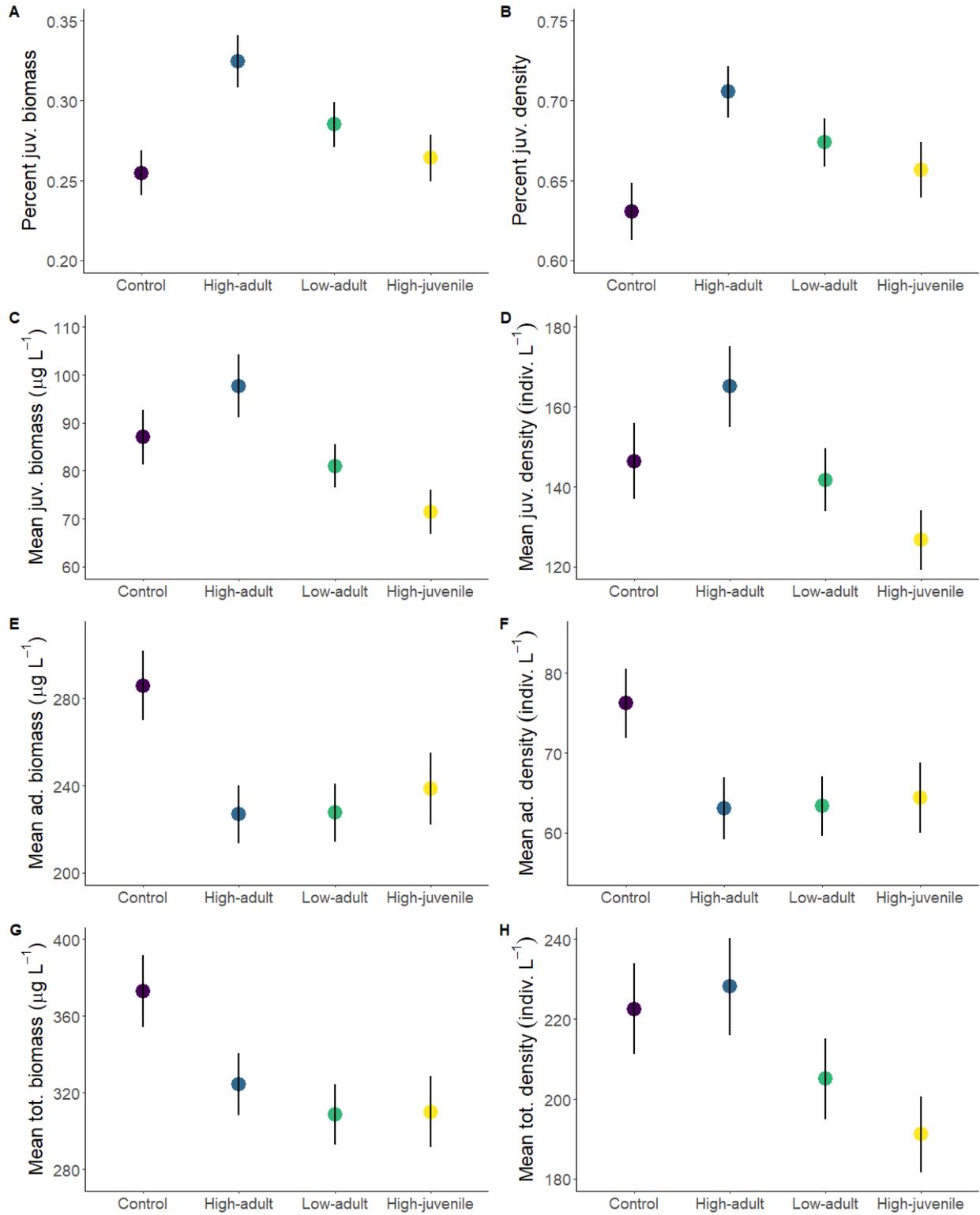
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592 **Fig. 1.** *Daphnia* biomass removed by harvesting treatments. (A) Time-series of mean juvenile  
 593 and adult biomass removed (averaged across treatment replicates) over the duration of the  
 594 experiment. Vertical black lines indicate the days on which *Daphnia* were harvested. (B) Mean  
 595 juvenile biomass removed per harvesting event  $\pm 1$  standard error over the duration of the  
 596 experiment (n = 11 harvesting events total). (C) Mean adult biomass removed per harvesting  
 597 event  $\pm 1$  standard error over the duration of the experiment.



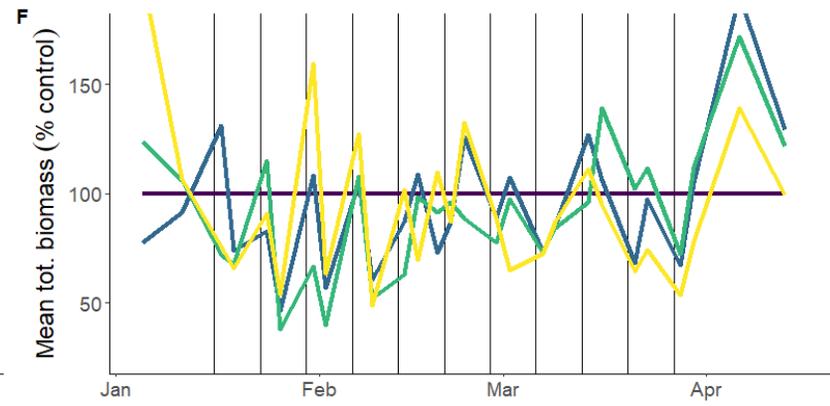
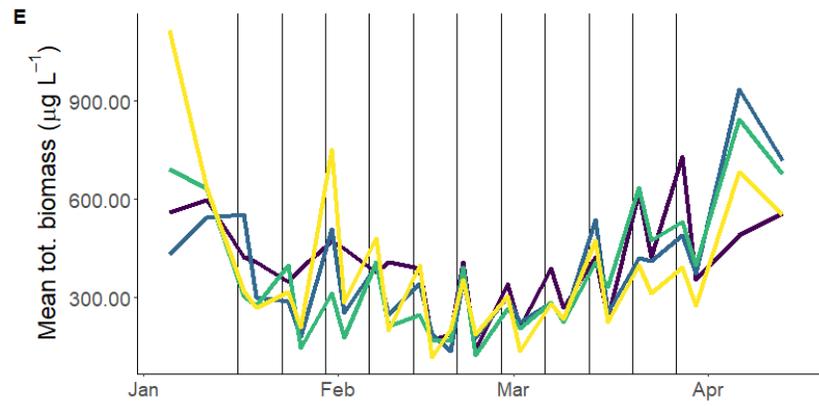
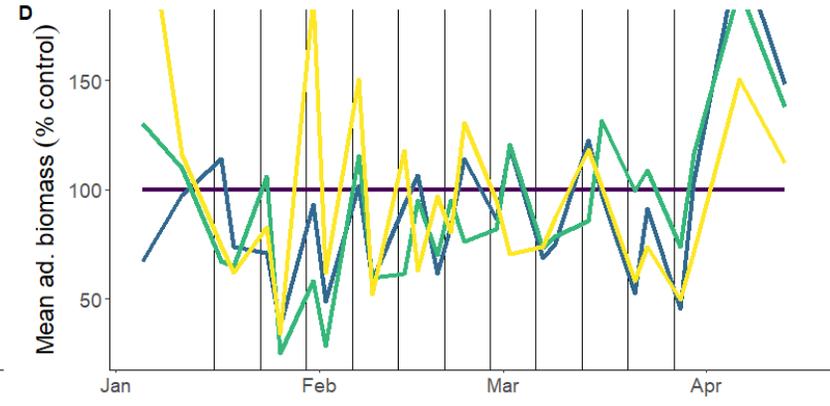
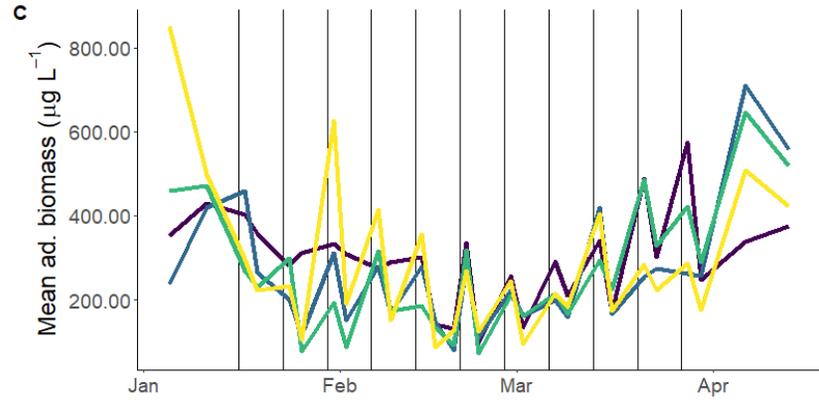
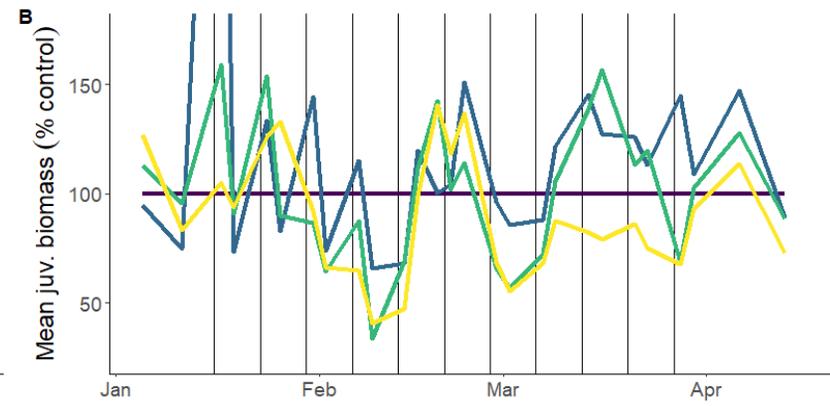
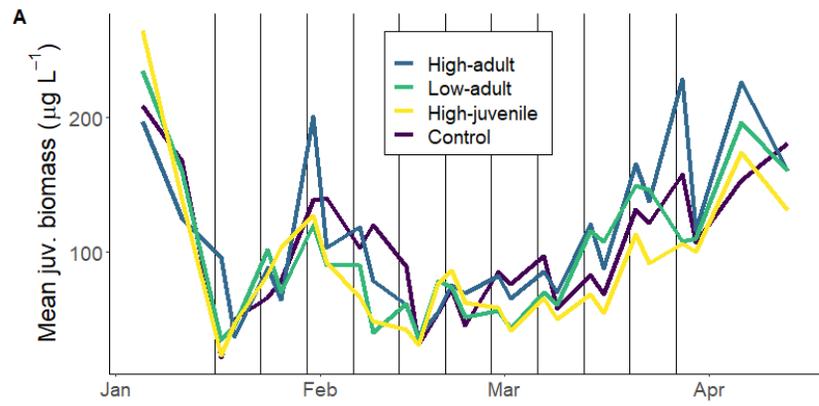
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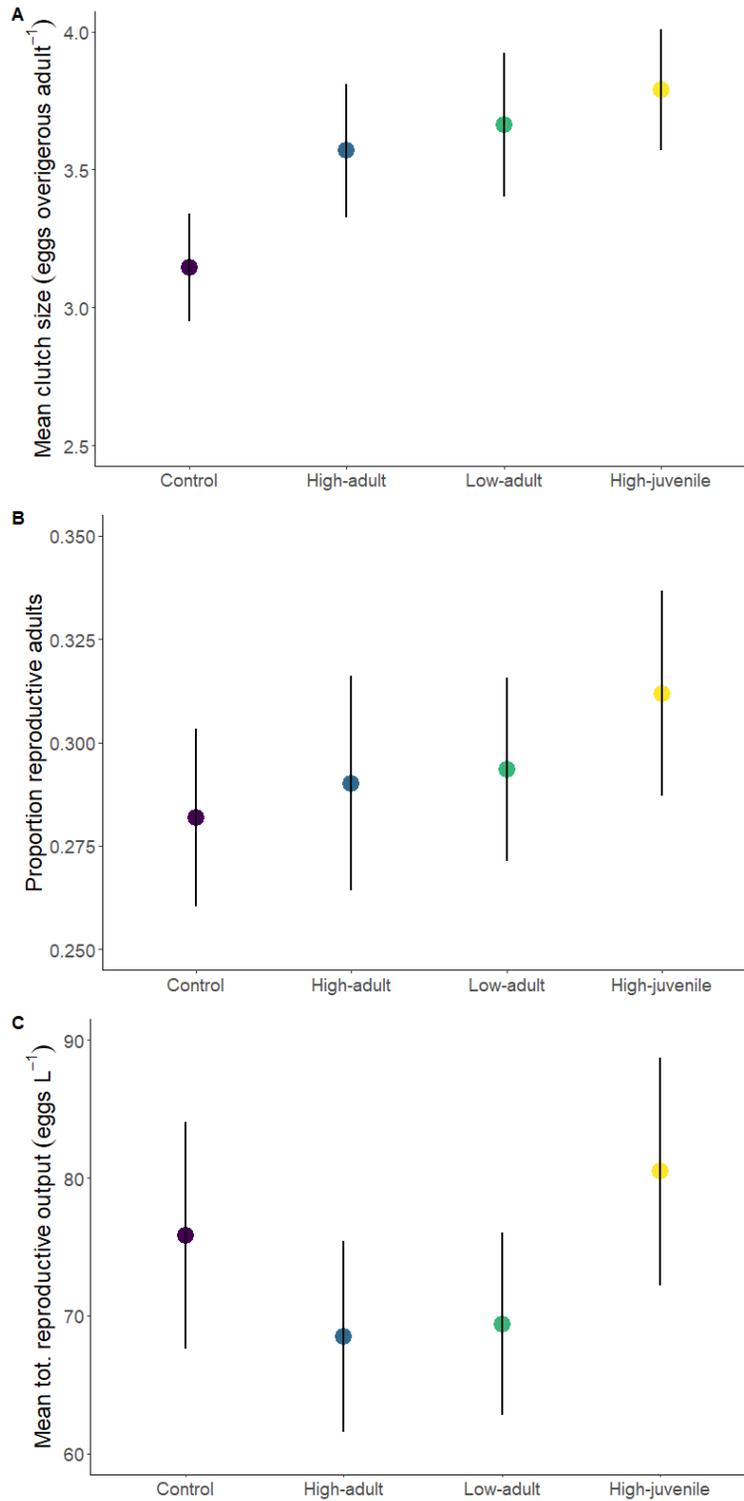
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**Fig. 2.** *Daphnia* population metrics (mean  $\pm$  1 standard error) averaged over the duration of the experiment for control and harvesting treatments. (A) Mean percent juvenile biomass. (B) Mean

601 percent juvenile density. (C) Mean juvenile biomass. (D) Mean juvenile density. (E) Mean adult  
602 biomass. (F) Mean adult density. (G) Mean total biomass. (H) Mean total density.



**Fig. 3.** Time-series of mean *Daphnia* biomass for control and harvesting treatments (A, C, E) and mean *Daphnia* biomass normalized as percent of control biomass (B, D, F) over the duration of the experiment. Vertical black lines indicate the days on which *Daphnia* were harvested. Error bars omitted for clarity.



**Fig. 4.** *Daphnia* reproductive responses to harvesting treatments averaged over the duration of the experiment. (A) Mean clutch size  $\pm$  1 standard error (a *per capita* measure). (B) Mean

proportion of reproductive (ovigerous) adults  $\pm$  1 standard error (C) Mean total reproductive output  $\pm$  1 standard error (a population-level measure). Total reproductive output was calculated by multiplying mean clutch size  $\times$  the proportion of ovigerous adults per sample  $\times$  total adult density per sample.