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Does masting scale with plant size? High reproductive variability and low synchrony in small and unproductive individuals

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Does masting scale with plant size? High reproductive variability and low synchrony in small and unproductive individuals

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Abstract

Background and Aims In a range of plant species, the distribution of individual mean fecundity is skewed and dominated by a few highly fecund individuals. Larger plants produce greater seed crops, but the exact nature of the relationship between size and reproductive patterns is poorly understood. This is especially clear in plants that reproduce by exhibiting synchronized quasi-periodic variation in fruit production, a process called masting.

Methods We investigated covariation of plant size and fecundity with individual-plant-level masting patterns and seed predation in 12 mast-seeding species: *Pinus pinea*, *Astragalus scaphoides*, *Sorbus aucuparia*, *Quercus ilex*, *Q. humilis*, *Q. rubra*, *Q. alba*, *Q. montana*, *Chionochloa pallens*, *C. macra*, *Celmisia lyallii* and *Phormium tenax*.

Key Results Fecundity was non-linearly related to masting patterns. Small and unproductive plants frequently failed to produce any seeds, which elevated their annual variation and decreased synchrony. Above a low fecundity threshold, plants had similar variability and synchrony, regardless of their size and productivity.

Conclusions Our study shows that within-species variation in masting patterns is correlated with variation in fecundity, which in turn is related to plant size. Low synchrony of low-fertility plants shows that the failure years were idiosyncratic to each small plant, which in turn implies that the small plants fail to reproduce because of plant-specific factors (e.g. internal resource limits). Thus, the behaviour of these sub-producers is apparently the result of trade-offs in resource allocation and environmental limits with which the small plants cannot cope. Plant size and especially fecundity and propensity for mast failure years play a major role in determining the variability and synchrony of reproduction in plants.

Keywords: Fecundity, mast seeding, plant reproduction, predator satiation, seed predation, super-producers.

Introduction

Recruitment of the majority of plant species is limited by the availability of seeds (Clark *et al.*, 2007). Thus, individual variation in fecundity within plant populations is a life-history parameter of high evolutionary and ecological significance (Herrera and Jovani, 2010; Moran and Clark, 2012). In a range of plant species, variation among individuals in fecundity tends to be high, with seed production dominated by a few highly fecund individuals (Greenberg, 2000; Herrera and Jovani, 2010; Pesendorfer *et al.*, 2016). Older and larger plants generally produce greater seed crops, but the exact nature of this relationship between plant size and reproduction is poorly understood (Thomas, 2011; Hossain *et al.*, 2017; Pesendorfer *et al.*, 2020). This knowledge gap is especially clear in perennial plants that reproduce through masting cycles, characterized

by large, synchronized annual variation in fruit production (Kelly, 1994; Vacchiano *et al.*, 2018).

Recent studies imply that large within-species differences in the extent of the inter-annual variability and synchrony may be driven by the age or size of individual plants, yet they report contrasting relationships (Minor and Kobe, 2017; Pesendorfer *et al.*, 2020). On one hand, older (and presumably larger) European temperate forest trees are more fecund and more variable (Pesendorfer *et al.*, 2020). On the other hand, larger (and presumably older) North American temperate trees produced more seeds, and the top 10 % of the most fecund individuals (called ‘super-producers’) had lower annual variation of seed production (Minor and Kobe, 2017). The size-dependent differences in masting can amplify or reduce the fitness differences among individuals varying in fecundity, as fitness of masting plants depends on the functional benefits that the inter-annual variability and synchrony provide.

The two functional benefits of masting with the most frequent empirical evidence are predator satiation and increased pollination efficiency; here, we focus on the first (Pearse *et al.*, 2016). During conditions of predator satiation, large variation in crop size causes seed predators to starve in low-seed years and to experience satiation in mast years (Satake and Bjørnstad, 2004; Linhart *et al.*, 2014). Even though predator satiation depends on population-level patterns of reproduction, individual plants gain fitness benefits according to their individual degree of annual variability and synchronization of reproduction (Ims, 1990; Koenig *et al.*, 2003; Satake and Bjørnstad, 2004; Żywiec *et al.*, 2013). Seed predation by specialist insect granivores could be especially susceptible to plant-level changes in annual variability or synchrony because of their relatively low mobility (Koenig *et al.*, 2003; Bogdziewicz *et al.*, 2020). Consequently, if large and fecund plants produce seeds more regularly, they could experience increased seed losses if this stable seed supply results in higher local average survival of insect cohorts and a localized build-up of insect populations (Maeto and Ozaki, 2003; Higaki, 2016; Bogdziewicz *et al.*, 2017).

Here, we investigate the covariation of fecundity with respect to plant size, masting patterns and pre-dispersal seed predation using long-term (12–30 years) data for a diverse set of 12 species. We define fecundity as the mean seed production of an individual plant. (1) First, we tested whether fecundity correlates with plant size (Greenberg, 2000; Minor

and Kobe, 2017). Next, we tested the relationships among overall fecundity, variability and synchrony. Our hypothesis was that inter-annual reproductive variation and synchrony are linked (correlated) primarily to fecundity by the frequency of non-reproductive years (failure years). Specifically, we predicted that (2) small plants with low fecundity would experience reproductive failure more frequently than larger ones, (3) individual plant variability would be elevated by the proportion of failure years in time series, and (4) synchrony would be reduced by the proportion of size-driven failure years. Consequently, if all our predictions held, (5) reproductive variation across years would be higher and (6) synchrony would be lower in small and unproductive individuals. As we predicted that high-fecundity plants will produce seeds more regularly in comparison with low-fecundity individuals, we predicted that (7) seed predation correlates positively with fecundity. In addition to analysing correlations between size, fecundity and masting patterns as continuous variables, we also characterized seed production patterns for super-producers (10 % most fecund plants) versus the remainder in each species, following Minor and Kobe (2017). This categorical analysis was done to contrast the results of a dichotomous versus continuous approach.

Materials and methods

Study species

Annual variation in reproductive effort was monitored for individual plants by collection of all the cones on the whole plant (*Pinus pinea*), counting all inflorescences and seed pods (*Astragalus scaphoides*), counting all fruits (*Sorbus aucuparia*), counting fruits on selected branches (*Quercus ilex*, *Q. humilis*), using seed traps (*Q. rubra*, *Q. alba*, *Q. montana*) or counting all flowers (*Chionochloa pallens*, *C. macra*, *Celmisia lyallii* and *Phormium tenax*). For ten species (*S. aucuparia*, *P. pinea*, *Q. ilex*, *Q. humilis*, *Q. rubra*, *Q. alba*, *Q. montana*, *C. pallens*, *C. macra*, *C. lyallii*) we also collected plant size data, while for another subset of seven species we collected pre-dispersal seed predation data by insects (*S. aucuparia*, *P. pinea*, *Q. ilex*, *Q. humilis*, *Q. rubra*, *Q. alba* and *Q. montana*). All 12 species show clear masting behaviour (Kelly *et al.*, 2000, 2013; Espelta

Table 1. Summary the ecology of study species, sites and field procedures. Details are given in Supplementary Data Appendix S1

<i>Species</i>	<i>Location</i>	<i>Number of sites</i>	<i>Individuals monitored</i>	<i>Life form</i>	<i>Reproductive trait measured</i>	<i>Collecting method</i>	<i>Plant size measurement</i>
<i>P. pinea</i>	Spain	52	187	Tree	Cones	Census	DBH
<i>A. scaphoides</i>	USA	4	507	Herb	Inflorescences	Census	NA
<i>S. aucuparia</i>	Poland	1	299	Tree	Fruits	Census	DBH
<i>Q. ilex</i>	Spain	17	225	Tree	Acorns	Count on selected branches	DBH
<i>Q. humilis</i>	Spain	17	172	Tree	Acorns	Count on selected branches	DBH
<i>Q. rubra</i>	USA	3	44	Tree	Acorns	Seed trap	DBH
<i>Q. alba</i>	USA	3	51	Tree	Acorns	Seed trap	DBH
<i>Q. montana</i>	USA	2	33	Tree	Acorns	Seed trap	DBH
<i>C. pallens</i>	New Zealand	5	217	Grass	Inflorescences	Census	BA
<i>C. macra</i>	New Zealand	5	125	Grass	Inflorescences	Census	BA
<i>C. lyallii</i>	New Zealand	3	94	Herb	Inflorescences	Census	Rosettes
<i>P. tenax</i>	New Zealand	1	37	Herb	Inflorescences	Census	NA

DBH, diameter at breast height; BA, basal area; NA, data not available.

et al., 2008; Crone *et al.*, 2009; Calama *et al.*, 2017; Bogdziewicz *et al.*, 2018a, 2019). Furthermore, in ten of the 12 focal species, our past investigations indicated that masting decreases the proportion of seeds attacked by pre-dispersal seed predators in high-seed years (Kelly and Sullivan, 1997; Kelly *et al.*, 2000; Crone and Lesica, 2004; Espelta *et al.*, 2008; Żywiec *et al.*, 2013; Calama *et al.*, 2017). The exceptions are two North American oaks, *Q. alba* and *Q. montana* (Bogdziewicz *et al.*, 2018). A description of the ecology of the study species, sites and field procedures is given in the Supplementary Data Appendix S1 and is summarized in **Table 1**.

Data analysis

Reproductive traits. For each plant, we described masting behaviour using two metrics commonly used in studies of mast seeding: coefficient of variation (CVi, calculated as the ratio between the standard deviation and the mean) and amongindividual synchrony (S) (Koenig *et al.*, 2003; Crone *et al.*, 2011). The synchrony of seed production (technically diaspores, or of fruits/flowers/cones, referred to as ‘seeds’ throughout the

text for convenience) of each individual was measured by the average Pearson pairwise correlation of seed production of an individual plant with all other individuals observed at the site (Koenig *et al.*, 2003). We also estimated the proportion of failure years, calculated as the ratio of years for which no seeds were recorded for an individual plant versus the number of years that plant was monitored. To avoid bias due to limited sampling, we used only individuals that were monitored for at least 10 years.

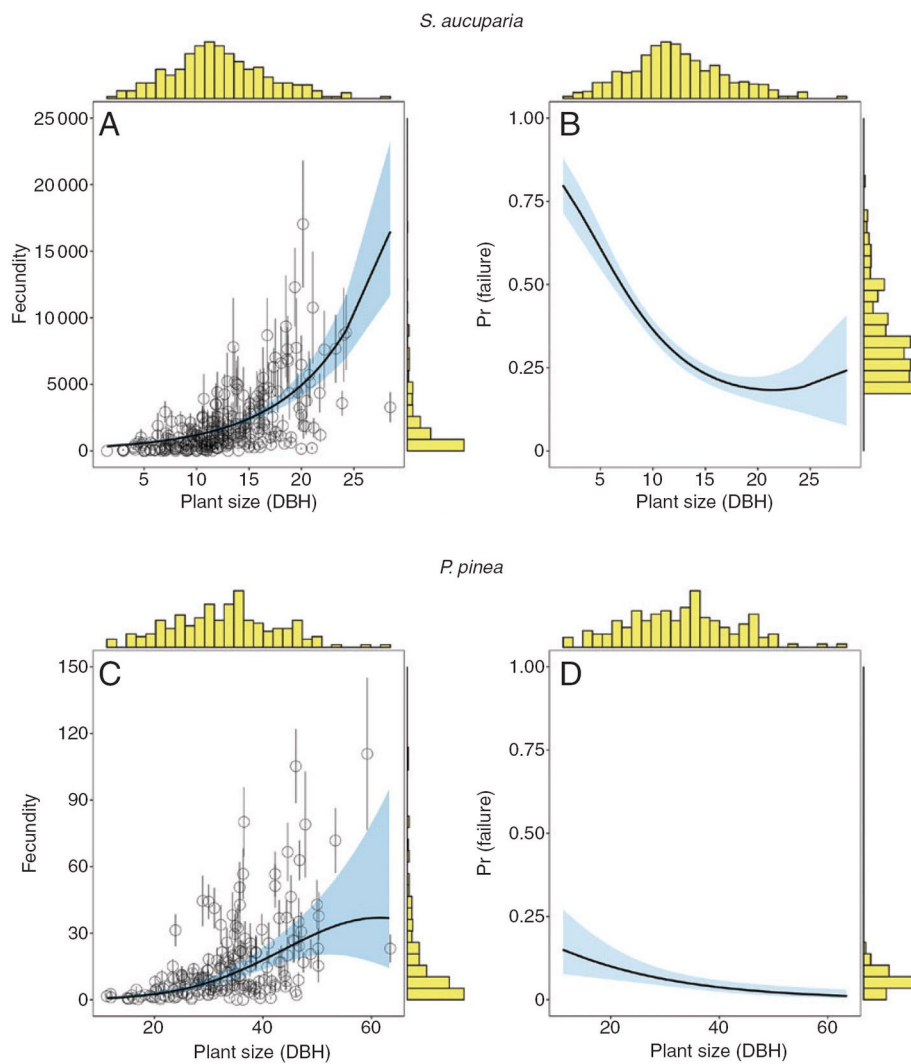
In addition, to contrast the results of the dichotomous versus continuous approaches, we also defined ‘super-producers’ as the subset of each population that was above the 90th percentile of individual-plant fecundity (Minor and Kobe, 2017). Fecundity was calculated as the annual mean reproductive effort by each plant (total number of seeds produced by an individual plant divided by the number of years a plant was monitored), to adjust for the differences in the number of sampling years among individuals. We then characterized seed production patterns for each sub-population (super-producers versus the other individuals) in each species, using the masting metrics (CVi and S).

Fecundity versus other reproductive traits. To explore whether variation in fecundity was related to focal plant size (prediction 1), we used zero-inflated negative binomial (ZINB) mixed models with plant size (diameter at breast height or basal area index, depending on the species; see Supplementary Data Appendix S1) included as the predictor in both the negative binomial and binomial part of the model. We used annual seed counts as the response. For each species, we fitted four candidate models, including plant size as a linear or quadratic term in all possible combinations in both logistic and negative parts of the model. Study site (with the exception of those for *S. aucuparia* and *P. tenax*, which were monitored on only one site) and tree ID were included as random terms. To account for differences in sampling effort at the plant level, each model included the log-transformed number of sampling years as an offset. The best model was chosen based on the standard Akaike information criterion and only that one is reported. In the case of *P. pinea* and *Q. montana*, due to ZINB model convergence issues, we first modelled annual seed production as a function of size using negative binomial mixed models and then, using binomial mixed models, regressed probability of failure (no reproduction in a particular year) with plant size.

Next, to explore the putative link between fecundity and masting patterns (predictions 2–4), we built generalized linear mixed models (GLMMs) that included (1) the proportion of failure years as a response and tree-level mean fecundity as a predictor (prediction 2), (2) CV_i as the response and proportion of failure years as a predictor (prediction 3), and (3) synchrony as the response and proportion of failure years as the predictor (prediction 4). The first model used a binomial family error term and logit link (prediction 2), and the second one (prediction 3) used a Gaussian error term and identity link. To analyse associations between synchrony and proportion of failure years (prediction 4), we used GLMMs with Tweedie distribution and logit link, with the response normalized as $y_i = (y_i + 1)/2$. In the next step, we correlated reproductive variation (prediction 5) and synchrony (prediction 6) with tree-level mean fecundity using GLMMs with Gaussian error term and identity link or Tweedie distribution and logit link, respectively. Finally, we used a GLMM with a binomial family error term and logit link to test whether the annual proportion of damaged seeds correlates with tree-level mean fecundity (prediction 7). In all analyses, we built separate models for each species and considered both linear and quadratic effects of an explanatory term. In addition, when testing predictions 2 and 5 we fitted the relationship between the response and predictor as a self-starting asymptotic function. In all models, we used site ID as a random effect (with the exception of *S. aucuparia* and *P. tenax*, which were monitored on only one site). In the GLMMs testing prediction 7 (predation versus tree-level mean fecundity) we used also tree ID as a random effect and included temporal autocorrelation (lag1). We calculated marginal effects (i.e. the proportion of variance explained by fixed effects) and conditional effects (i.e. the proportion of variance explained by fixed and random effects) with R^2 for our models (Nakagawa and Schielzeth, 2013). All statistics were run in R version 3.4.1 (R Core Team, 2018). We used the glmmTMB package (Brooks *et al.*, 2017) to fit all the models and the DHARMA (Hartig, 2017) package to validate them based on visual inspection of residual patterns.

Results

In nine of ten species for which we had plant size data (all except *Q. montana*), fecundity was significantly related to plant size (Supplementary Data Table S1, **Fig. 1**). Moreover, in seven out of ten species (exceptions were *C. lyallii*, *Q. montana* and *Q. alba*), larger plants experienced fewer failure years (Supplementary Data Table S1). The variation explained by these models ranged from 5 to 59 % across all species. Similarly, in all species, more fecund plants experienced fewer failure years, and fecundity explained 6–80 % (mean = 22 %) of variance in reproductive failures (Supplementary Data Table S2).



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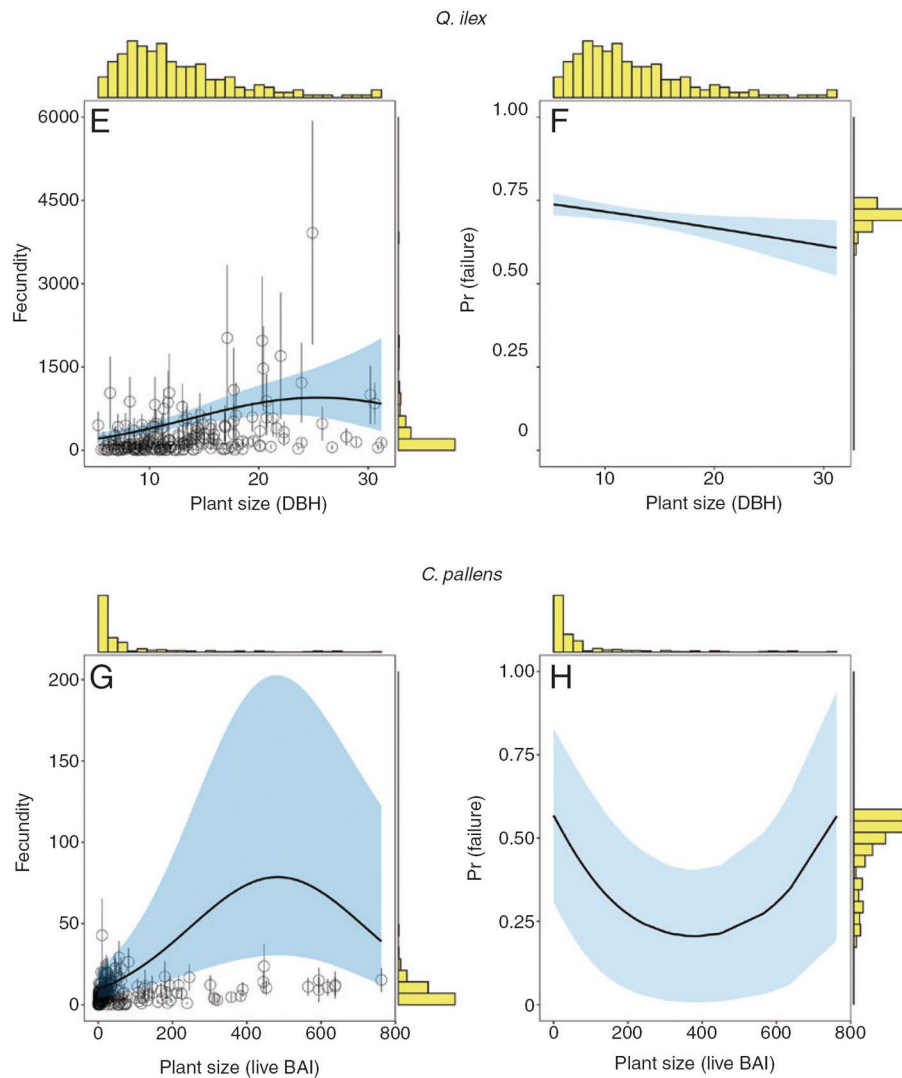


Fig. 1. Relationships between fecundity (modelled as annual seed counts; A, C, E, G) or probability of failure [Pr(failure)] (B, D, F, H) and plant size. Solid lines show statistically significant ZINB model predictions and associated 95 % confidence bands. Points in panels (A), (C), (E) and (G) show long-term mean fecundity (\pm s.e.) of individual plants. This figure presents a subset of the studied species (see Supplementary Data Table S1 for results for all species). The models included tree ID and site as random effects (see Materials and methods for details). DBH, diameter at breast height; BAI, basal area index.

Reproductive variability as measured by CVi was strongly related to the proportion of failure years in the time series of all species, and failures explained most of the variance (mean = 75 %) (Supplementary Data Table S3, **Fig. 2**). Similarly, reproductive synchrony was strongly related to the proportion of failure years, and failures explained large portions of the variance (mean = 44 %) (Supplementary Data Table S4, **Fig. 2**). This relationship was significant in 11 out of 12 species tested (all but *Q. humilis*). Consequently, variability (CVi) was negatively, non-linearly related to fecundity in all species (Supplementary Data Table S5), and fecundity explained, on average across species, 41 % of the variance in CVi. Importantly, inter-annual variability was stable across most of the fecundity range and increased sharply for the lowest-fecundity individuals (**Fig. 3**). In addition, fecundity was significantly positively correlated to synchrony in ten out of 12 species (all but *Q. montana* and *Q. humilis*), and explained a significant portion of the variance (mean = 49 %, Supplementary Data Table S6, **Fig. 3**). Seed predation was positively related to fecundity in only two species: *Q. ilex* and *Q. humilis* (Supplementary Data Table S7, **Fig. 4**).

In the dichotomous comparisons of the super-producers (10 % most fecund plants) versus the remainder of individuals in each species, these highly fecund plants had lower CVi and higher synchrony (S). Depending on the species, superproducers produced 14–53 % of the total seed count in the population (Supplementary Data Table S8).

Discussion

The sources of within-species variation in masting behaviour are largely unknown, but our study shows that significant portions of this variation are driven by differences in mean fecundity, which in turn is correlated with plant size. Small plants produce few seeds, and fail to produce seeds frequently, which elevates their reproductive variation across years and sharply decreases synchrony with other individuals in the population. This result partially agrees with the past observation that larger trees produce more seeds with lower inter-annual variability (Minor and Kobe, 2017). Specifically, a dichotomous comparison of the top 10 % most fecund plants with the remaining 90 % implies that super-producers tend to have lower annual variability of seeding and greater synchrony (Supplementary Data Table S8). Greater synchrony was hypothesized to give super-producers the ability to reap the benefits of

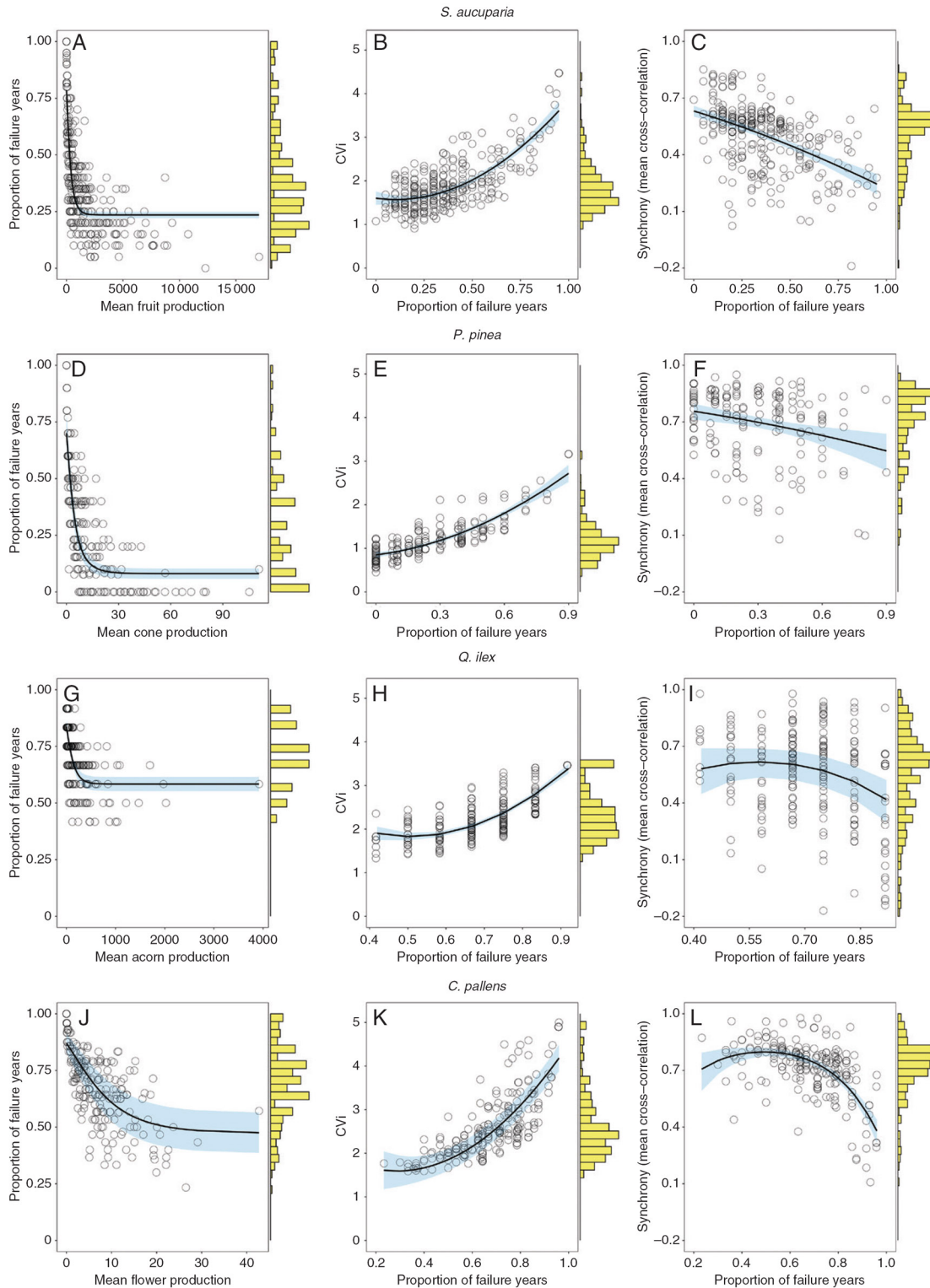


Fig. 2. Relationships between proportion of years with reproductive failure (no reproduction observed) of a plant, plant-level CVI in annual reproduction, synchrony and long-term mean plant fecundity (overall number of fruits produced/number of years a plant was monitored). Solid lines show statistically significant GLMM predictions and shaded areas show associated 95 % confidence bands. This figure presents a subset of the studied species (see Supplementary Data Tables S2–4 for results for all species). The models included site as a random effect (see Materials and methods for details).

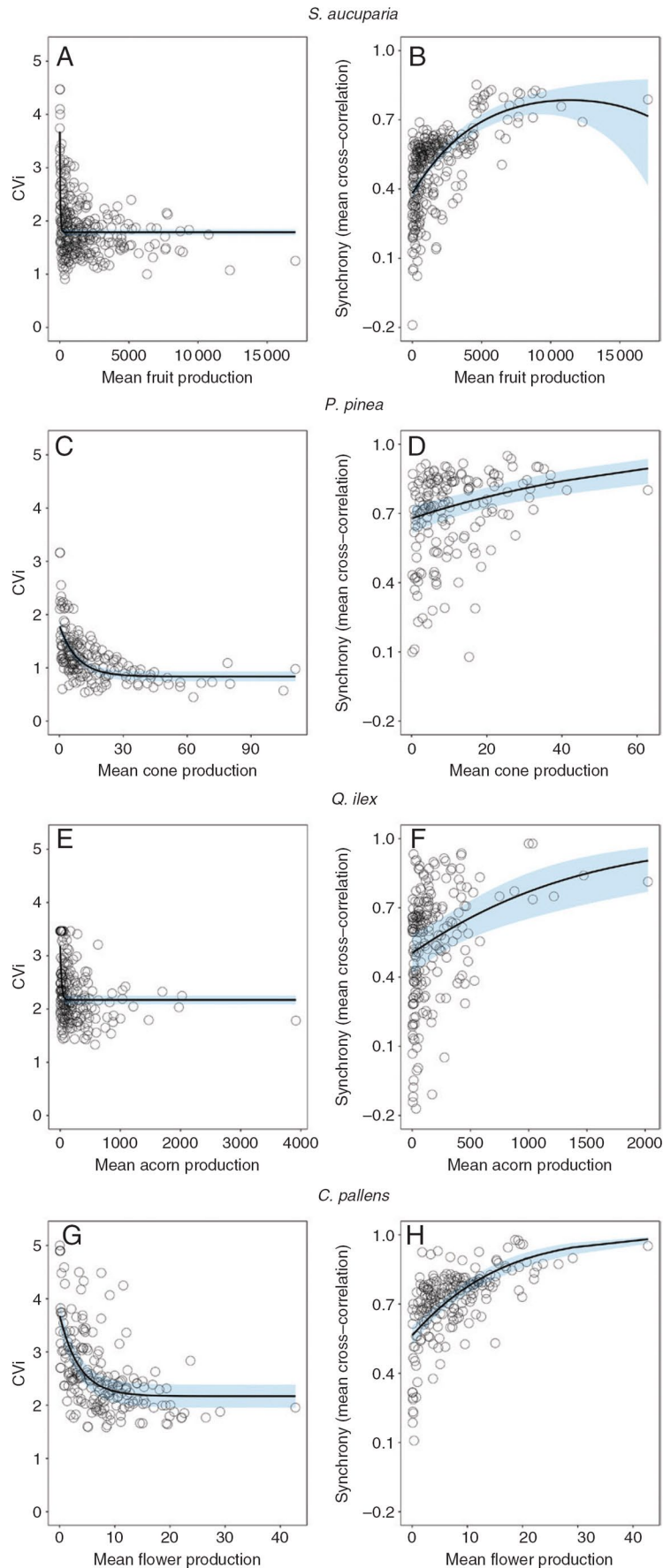


Fig. 3. Scatterplots of plant-level measures of CVi in annual reproduction, synchrony and long-term mean fecundity (overall number of fruits produced/ number of years a plant was monitored). Solid lines show statistically significant GLMM predictions and associated 95 % confidence bands. This figure presents a subset of the studied species (see Supplementary Data Tables S5–6 for results for all species). The models included site as a random effect (see Materials and methods for details).

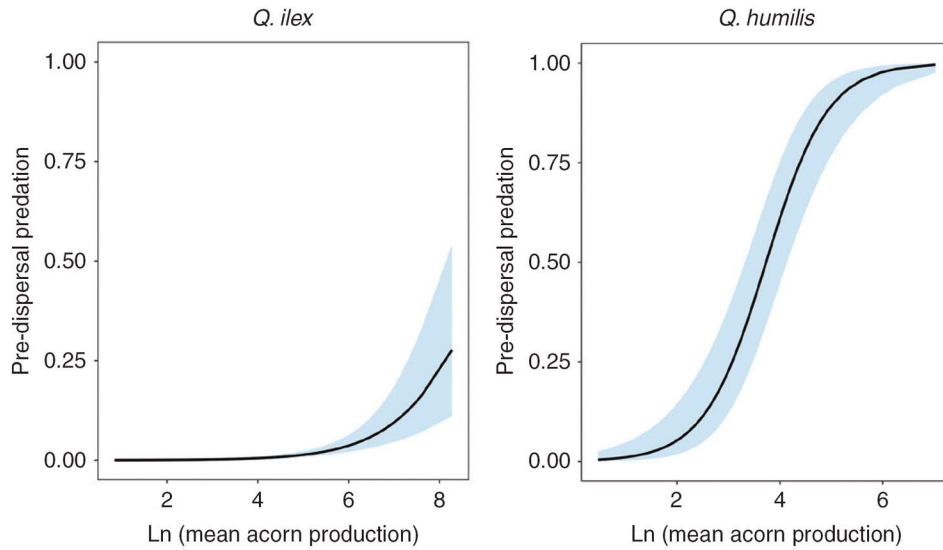


Fig. 4. Plant-level pre-dispersal predation rate versus fecundity (overall number of fruits produced/number of years a plant was monitored). Solid lines show statistically significant GLMM model predictions and associated 95 % confidence bands. This figure presents a subset of the studied species in which the relationship was significant (see Supplementary Data Table S7 for results for all species). The models included tree ID and site as random effects (see Materials and methods for details).

masting while also governing regular seed production over time (Minor and Kobe, 2017). Our comparison of this categorical approach with a continuous one indicates that the notion that super-producers behave differently is driven by the smallest and least fertile plants in the population. These plants drive the mean up (in the case of inter-annual variability) or down (in the case of synchrony) for all non-super-producers. In other words, in terms of masting patterns, there are no super-producers, but rather normal plants and sub-producers. This distinction is important biologically, as it shows that rather than the most fecund plants behaving differently from other plants in the population, the least fertile individuals are the outliers.

As well as the importance of continuous rather than dichotomous analyses, our analyses suggest a few other important methodological lessons in the study of individual plant variation. High CV values occur in individuals and populations that have many years with zero reproduction. This is not surprising given that CV is directly mathematically linked to occurrence of zeros in the time series (Crone *et al.*, 2011). In time series with many years with failure, the CV is much less influenced by the amount of reproduction in non-zero years (Crone *et al.*, 2011).

Similarly, as in the case of CV, among-plant variation in synchrony was also large and linked to the occurrence of zero years. Shared failure years (e.g. those in response to environmental disturbances like drought, cf. Rees *et al.*, 2002; Bogdziewicz *et al.*, 2018b) would give small and infertile plants greater synchrony, but in fact small plants have lower synchrony. Thus, the failure years were idiosyncratic to each small plant, which implies that these plants fail to reproduce because of plant-specific factors (e.g. internal resource limits), combined with selection for small plants to allocate fewer resources to reproduction and more to growth until they are larger or taller (Miller *et al.*, 2008; Thomas, 2011). Therefore, the failures of the least fertile plants could be a result of resource allocation trade-offs and environmental limits with which the small plants cannot yet cope. In addition, many plants shift resource allocation from growth to reproduction with increasing plant maturity (Thomas, 2011). Understanding the relationship between size and age is not possible in this study because we did not have estimates of plant age for all individuals, noting especially that our multispecies data include many herbaceous plants. Evaluating effects of age *per se*, as well as other differences among individual plants within size classes, could be an interesting area for future research.

In five out of seven species for which we had seed predation data, higher seed predation was not associated with higher fecundity, even though more fecund plants reproduced more regularly. This pattern suggests that failure years do not have a decisive influence on the insect seed predators' populations in these species, possibly because predators are able to move between plants. In support of this idea, recent studies show greater insect immigration into asynchronous trees when the population-level seed production is low (Bogdziewicz *et al.*, 2018a). In two Mediterranean oaks in which we detected a positive correlation between seed predation and fecundity, infrequent failures likely lead to lower insect emigration or higher survival, which over the long term leads to a build-up of the insect populations (Bogdziewicz *et al.*, 2017; Pérez-Ramos *et al.*, 2017). We hypothesize that the species-specific relationship between regular seeding and pre-dispersal seed predation we detected is caused by interactive effects of plant and predator population traits, including predators' costs of migration in low-seed years, insect mobility, and the density of the plant population (Moreira *et al.*, 2017; Bogdziewicz *et al.*, 2018c). This warrants further investigation, but will require even larger datasets.

In closing, our analysis shows that, presumably because they are under constant selection to gain economies of scale, within a species plants over a wide size range have remarkably similar masting strategies. The only departures from this are the least fecund plants (sub-producers), which might differ because of fundamental constraints on the smallest feasible reproductive output. Small and low-fertility plants often fail to produce seeds, which elevates their inter-annual variability and decreases reproductive synchrony. This pattern was consistent among all studied species, which included both angiosperms and gymnosperms, evergreen and winter-deciduous species, trees and herbs. A careful generalization based on the sample of 12 species studied here implies that the individual-level differences in annual reproductive variation can be associated with plant size and the propensity of small plants to shift resource allocation away from reproduction in favour of growth. Plant size and fecundity play a major role in determining the variability and synchrony of reproduction in plants. Understanding the mechanisms of the size–fecundity–synchrony relationship could be an important next step in predicting how mast seeding will change in changing environments.

Supplementary data

Supplementary data follow the **Literature cited** section and consist of the following.

- Appendix S1: study species and data collection.
- Table S1: summary of ZINB mixed models regressing variation in fecundity with plant size.
- Table S2: summary of generalized mixed models regressing the variation in the proportion of failure years a plant experienced during the study period versus that plant's mean fecundity.
- Table S3: summary of linear mixed models regressing the variation in annual variability of reproduction of a focal plant with the proportion of failure years in the time series.
- Table S4: summary of generalized mixed models regressing the variation in synchrony of reproduction of a focal plant with the proportion of failure years in the time series.
- Table S5: summary of linear mixed models regressing the variation in plant CVi versus the plant's fecundity.
- Table S6: summary of generalized mixed models regressing the variation in plant reproductive synchrony versus the plant's fecundity.
- Table S7: summary of generalized mixed models regressing the variation in pre-dispersal seed predation versus log-transformed mean plant fecundity.
- Table S8: annual variability and synchrony of seed production in the studied species.

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Appendix S1.

Does mast seeding scale with plant size? Large reproductive variability and low synchrony in small and unproductive plants

Study species and data collection

Pinus pinea

Pinus pinea (Pinaceae) lives up to 200 years. The fruiting process covers a four-year period between reproductive bud differentiation, cone opening and subsequent seed dispersal. The study sites were located in Central Range of Spain, at a mean altitude of 600 - 900 m. The climate of the study area is genuine Mediterranean, defined by a long, hot, dry summer period and humid winters and autumns. Average annual rainfall in the area is 648 mm. Average annual temperature is 13.7 °C.

Cone data was collected over 52 sample plots installed in even-aged stone pine stands. The plots are circular, of variable size, and include 20 trees. Plots were installed aiming to cover the whole range of site, age and stocking conditions within the territory. All the cones from the five trees nearest to the centre of the plot were manually harvested by specialized climbers every autumn. 260 trees were sampled in total. The cones cropped from each tree were classified as 'healthy' or 'damaged'. In Central Spain, the most common damage is caused by *Dioryctria mendacella*, which is in accordance with our observations, where more than 95% of the cones classified as damaged were infested by this species.

Astragalus scaphoides

Astragalus scaphoides (Fabaceae) is an iteroparous legume that lives ~20 years, endemic to high-elevation sagebrush steppe in a small area of Beaverhead County in southwestern Montana and adjacent Lemhi County in east-central Idaho USA. The climate is semi-arid; mean annual precipitation in Lemhi and Beaverhead counties is about 250 mm, with peak rainfall in May. *A. scaphoides* does not reproduce vegetatively, and is visited by a number of generalist bumblebees (*Bombus* spp.) and solitary bees. Flowering occurs from late May to mid-June. In most years, plants dehisce seeds by mid-July.

Eight permanent monitoring transects were established at Sheep Corral Gulch, Montana (1987), Haynes Creek, Idaho (1987), and McDevitt Creek, Idaho (1989), and Reservoir Creek, Montana (2004) USA (two transects at each site). Each transect consisted of

50 adjacent 1-m² mapping quadrats placed along a transect line. In early July, when most fruits were mature and vegetative plants had not yet senesced, the position of each *A. scaphoides* plant encountered in the quadrats was mapped, and total inflorescence production as well as number of fruits produced were recorded.

Sorbus aucuparia

Sorbus aucuparia (Rosaceae) is a deciduous fleshy-fruited tree that lives 100–150 years. Flowering and fruit maturation occur within one year. The flowers are pollinated by a wide range of insects. Seed production data were collected in a subalpine old-growth spruce forest on Babia Góra massif (1,725 m a.s.l.) in the Western Carpathians (Poland) at an altitude of 1170–1310 m. The climate is cool with 3.3 °C mean annual temperature, 1,470 mm mean annual rainfall, and a snow-free period of 7 months.

We demarcated a 27-hectare (564 × 480 m) rectangular plot. All rowan trees in the plot with diameter at least 4 cm were mapped. The analysis used a group of 30 trees, i.e., a random subset of the population for which fruit predation was recorded. The whole plot is situated within one forest type and sandstone is the only bedrock type. Each year, all trees were searched for fruits at the beginning of September before birds began to feed on them. Fruit production was measured by binocular observations, counting the number of infructescences (corymbs with fruits) on individual trees. For each tree, in five infructescences all fruits were counted. The fruit production of a tree was determined as the product of its number of infructescences and the average number of fruits in five infructescences. To study pre-dispersal seed predation, three infructescences per tree (from different distant branches) were collected at the beginning of September. Ten fruits were taken randomly from each infructescence and dissected for seeds predated by apple fruit moth *Argyresthia conjugella*. Because there were no remnants of seeds in many fruits, it was not possible to estimate the number of seeds predated. For this reason, we took the percentage of fruits with signs of *A. conjugella* feeding (infested fruits) as the measure of predation at the individual level.

Quercus ilex and *Q. humilis*

The evergreen *Quercus ilex* (section *Cerris*) and the winter-deciduous *Q. humilis* (section *Quercus*) (both Fagaceae) are two Mediterranean, wind-pollinated, long-lived (up to >1000 years) oaks. The two species differ in their leaf habit but they share other life-history traits, such as fruit development, pollination, and maturation of acorns in one year. Data on

their seed production were collected in Collserola Natural Park (Barcelona, NE Iberian Peninsula). Climate is Mediterranean, with 620 mm of mean annual precipitation and mean monthly temperatures with a maximum of 23.3°C in August and a minimum of 7.9°C in January.

We monitored acorn production from 1998 to 2009 in 17 sampling sites (255 trees, 15 per plot) in holm-oak-dominated forests. Trees were tagged and the number of branches per tree was estimated using a regression model between crown projection and number of branches previously constructed for a subsample of trees. We counted acorn production and number of infested acorns (e.g., having a gnaw mark or hole caused by insect predation) on four branches per tree at the peak of the acorn crop (i.e. September). Then we estimated the total number of acorns produced per tree by multiplying the mean acorn production per branch and the number of branches per tree.

Quercus rubra, *Q. alba*, and *Q. montana*

Quercus rubra, *Q. alba*, and *Q. montana* oaks are winter-deciduous. They belong to two different sections of the genus *Quercus* (Lobatae: *Q. rubra* and *Quercus*: *Q. alba* and *Q. montana* respectively), which differ in length of time between flower production and acorn maturation. *Quercus* oaks produce flowers in the spring, which are fertilized and develop into mature fruit in the same year as they were pollinated. Lobatae oaks produce flowers that are pollinated during the spring of 1 yr, but the fruit matures the following year.

We monitored acorn production over 17 yr at three study areas in eastern Pennsylvania, USA: two in Luzerne County (Steele and Venesky) and a third in Schuylkill County (Hawk Mountain). Red and white oaks were monitored at all three sites, while chestnut oaks were not present at one site (the Steele site). We monitored acorn production by individual oak trees by placing two seed traps under each of 15 individuals of each species at each site from 2001 to 2017. Throughout the analysis, crop size per tree per year is the summed acorn count from the two seed traps. Each year, acorns were collected from seed traps biweekly from early August until seed fall ceased in mid-November. Acorns were bagged separately according to the tree of origin, then transported to the laboratory and refrigerated (4°C) until further processing. In the laboratory, we weighed, measured (width and length) and assessed insect infestation for individual acorns from each tree of each species.

Chionochloa pallens, *Chionochloa macra*, and *Celmisia lyallii*

The *Chionochloa* species are long-lived alpine tussock grasses (bunchgrasses) about 30-90 cm wide and tall which produce 0-150 tall inflorescences (culms) per plant each year. Individual plants of both species were mapped at Mt Hutt, Canterbury, New Zealand in permanent plots at several altitudes: 1070 m (mainly *C. pallens*, 1990-2019), 1540 m (mainly *C. macra*, 1992-2019), 1620 m (1996-2019) and 1540 m (2006-2019). For this analysis, we used only plants which had at least 10 years of data, and where there had been no disturbance (e.g. by skifield trail construction). For each plant in each year, the total number of culms produced was noted. Plant size was recorded as the total live basal area, measuring the basal diameter of the tussock to determine basal area, multiplied by the visually estimated fraction of the plant which had live green tissue.

Celmisia lyallii is a long-lived herbaceous rosette-forming alpine daisy which spreads clonally to form patches connected by rhizomes. Each rosette is about 20 cm diameter and can produce one or more flower stalks about 20 cm tall carrying a single inflorescence. Discrete patches were mapped in the *Chionochloa* plots at 1620 m (1996-2019), 1540 m (1997-2019) and 1520 m (2006-2019). For each patch, patch size was measured as the number of rosettes within it. Total counts of all inflorescence production per patch was recorded every year.

Phormium tenax

This is a long-lived herbaceous monocot. Clumps consist of a group of large interconnected rosettes. Each rosette carries leaves up to 2.5 m long, and can produce a large flowering stalk 3 m tall bearing several hundred flowers. Data were collected at Courtenay on the Canterbury Plains, New Zealand. A roadside planting of 37 discrete patches was mapped and total flower stalk production counted on each patch 2002-2019. No measurement of patch size was taken.

Tables S1 – S8 follow.

Table S1. Summary of Zero-inflated Negative Binomial mixed models regressing the variation in fecundity with the plant size (**prediction 1**). Each model included annual seed counts as a response and plant size as a predictor in conditional and binomial part of the model. R2m is the marginal (i.e. the proportion of variance explained by fixed effects) and R2c is the conditional (i.e. the proportion of variance explained by fixed and random effects) R2. For each species, we fitted four candidate models, including plant size as a linear or quadratic term in all possible combinations in both conditional and binomial parts of the model. The best one was chosen based on the standard AIC criteria and only that one is reported. In case of *P. pinea* and *Q. montana*, due to ZINB model convergence issues, we first modeled fecundity as a function of size using negative-binomial mixed models and then, using binomial mixed models, we regressed probability of failure (zero fecundity) with plant size.

Species	Effect	β	SE	z	p	β	SE	z	p	R2m	R2c
		Conditional part				Binomial part					
<i>S. aucuparia</i>	size	0.14	0.01	13.32	< 0.001	-39.68	4.14	-9.59	< 0.001	0.14	0.27
	size ²					15.32	4.10	3.74	< 0.001		
<i>Q. ilex</i>	size	19.53	4.15	4.71	< 0.001	-0.02	0.01	-2.52	0.012	0.06	0.32
	size ²	-7.73	3.43	-2.08	0.038						
<i>Q. humilis</i>	size	0.07	0.01	5.54	< 0.001	-0.04	0.01	-2.93	0.003	0.09 [#]	
<i>Q. rubra</i>	size	-2.28	1.98	-1.15	0.25	-0.047	0.02	-2.35	0.019	0.04 [#]	
	size ²	-4.84	1.91	-2.54	0.01						
<i>Q. alba</i>	size	0.01	0.01	2.05	0.04	0.003	0.01	0.29	0.769	0.03 [#]	
<i>C. pallens</i>	size	36.19	5.70	6.35	< 0.001	-23.40	6.50	-3.60	< 0.001	0.12	0.34
	size ²	-16.64	4.56	-3.65	< 0.001	20.65	5.81	3.55	< 0.001		
<i>C. macra</i>	size	27.10	4.12	6.58	< 0.001	-8.61	3.77	-2.28	0.022	0.10	0.49
	size ²	-10.51	4.12	-2.56	0.011	10.78	3.87	2.79	0.005		
<i>C. lyallii</i>	size	27.14	5.79	4.69	< 0.001	-0.01	0.01	-1.79	0.074	0.24 [#]	
	size ²	-12.94	4.53	-2.86	0.004						
		Negative-binomial model				Binomial model					
<i>P. pinea</i>	size	37.04	4.03	9.19	< 0.001	-0.05	0.02	-3.46	< 0.001	0.29*	0.59*
	size ²	-8.55	3.15	-2.71	0.007					0.05 ⁺	0.38 ⁺
<i>Q. montana</i>	size	0.02	0.02	1.00	0.316	-0.01	0.01	-0.86	0.391	0.01*	0.17*
										0 ⁺	0.05 ⁺

* R2 of negative binomial model, ⁺R2 of binomial model.

[#] Conditional R2 could not be computed due to low variance of the random effect.

Table S2. Summary of generalized mixed models regressing the variation in proportion failure years a plant experienced during the study period vs that plant mean fecundity (**prediction 2**). R2m is the marginal (i.e. the proportion of variance explained by fixed effects) and R2c is the conditional (i.e. the proportion of variance explained by fixed and random effects) R2. For each species, we fitted three candidate models – one that included a linear term of tree-level mean fecundity as a predictor, a second that included both linear and quadratic term, and the third one that fitted the relationship between the response and predictor as a Self-Starting asymptotic function. The best one was chosen based on the standard AIC criteria and only that one is reported.

Response: proportion of failure years

Species	Predictor: fecundity	β	SE	<i>z</i> -statistic	<i>p</i> -value	R2m	R2c
<i>P. pinea</i>	asymptotic	5.67	0.43	13.13	< 0.001	0.21	0.29
<i>A. scaphoides</i>	asymptotic	1.52	0.09	16.52	< 0.001	0.12	0.13
<i>S. aucuparia</i>	asymptotic	4.61	0.19	24.46	< 0.001	0.80*	
<i>Q. ilex</i>	asymptotic	5.06	0.50	10.12	< 0.001	0.06 [#]	
<i>Q. humilis</i>	asymptotic	5.27	0.40	13.12	< 0.001	0.14	0.15
<i>Q. rubra</i>	asymptotic	5.55	0.67	8.33	< 0.001	0.17 [#]	
<i>Q. alba</i>	asymptotic	4.61	0.69	6.70	< 0.001	0.09	0.10
<i>Q. montana</i>	asymptotic	4.85	0.80	6.04	< 0.001	0.09 [#]	
<i>C. pallens</i>	asymptotic	4.96	0.30	16.41	< 0.001	0.09	0.11
<i>C. macra</i>	asymptotic	4.42	0.40	11.02	< 0.001	0.09	0.09
<i>C. lyallii</i>	asymptotic	7.44	0.97	7.70	< 0.001	0.16	0.16
<i>P. tenax</i>	asymptotic	5.56	1.84	3.02	0.003	0.59*	

* Plants were monitored at one site, thus a model without site as random effect was fitted for that species. The R-squared was calculated as Pearson's correlation coefficient between the observed and predicted values of the response.

[#] Conditional R2 could not be computed due to low variance of the random effect.

Table S3. Summary of linear mixed models regressing the variation in annual variability of reproduction of a focal plant (CVi) with the proportion of failure years in the time series (**prediction 3**). R2m is the marginal (i.e. the proportion of variance explained by fixed effects) and R2c is the conditional (i.e. the proportion of variance explained by fixed and random effects) R2. For each species, we fitted two candidate models – one that included a linear term, and second that included both linear and quadratic term of proportion of failure years as predictors. The best one was chosen based on the standard AIC criteria and only that one is reported.

Response: CVi

Species	Predictor: proportion of zeros	β	SE	<i>z</i>-statistic	<i>p</i>-value	R2m	R2c
<i>P. pinea</i>	linear	5.06	0.24	21.23	< 0.001	0.73	0.73
	quadratic	1.21	0.24	5.06	< 0.001		
<i>A. scaphoides</i>	linear	13.32	0.43	30.67	< 0.001	0.80	0.81
	quadratic	5.20	0.35	14.67	< 0.001		
<i>S. aucuparia</i>	linear	6.95	0.38	18.41	< 0.001	0.75*	
	quadratic	2.50	0.38	6.63	< 0.001		
<i>Q. ilex</i>	linear	6.73	0.32	21.08	< 0.001	0.70 [#]	
	quadratic	2.78	0.32	8.70	< 0.001		
<i>Q. humilis</i>	linear	8.10	0.32	25.62	< 0.001	0.81	0.81
	quadratic	2.31	0.32	7.29	< 0.001		
<i>Q. rubra</i>	linear	3.16	0.37	8.66	< 0.001	0.69	0.73
	quadratic	1.28	0.34	3.76	< 0.001		
<i>Q. alba</i>	linear	4.74	0.32	15.05	< 0.001	0.84 [#]	
	quadratic	2.10	0.32	6.67	< 0.001		
<i>Q. montana</i>	linear	3.43	0.33	10.26	< 0.001	0.80 [#]	
	quadratic	1.51	0.33	4.52	< 0.001		
<i>C. pallens</i>	linear	9.03	0.50	17.95	< 0.001	0.66	0.72
	quadratic	2.38	0.43	5.57	< 0.001		
<i>C. macra</i>	linear	5.29	0.36	14.86	< 0.001	0.70	0.73
	quadratic	3.29	0.37	8.92	< 0.001		
<i>C. lyallii</i>	linear	4.94	0.39	12.78	< 0.001	0.63	0.71
	quadratic	1.12	0.39	2.85	0.004		
<i>P. tenax</i>	linear	0.66	0.08	7.92	< 0.001	0.84*	
	quadratic	0.33	0.08	3.98	< 0.001		

* Plants were monitored at one site, thus a model without site as random effect was fitted for that species. The R-squared was calculated as Pearson's correlation coefficient between the observed and predicted values of the response.

[#] Conditional R2 could not be computed due to low variance of the random effect.

Table S4. Summary of generalized mixed models regressing the variation in synchrony of reproduction of a focal plant (S) with the proportion of failure years in the time series (**prediction 4**). R2m is the marginal (i.e. the proportion of variance explained by fixed effects) and R2c is the conditional (i.e. the proportion of variance explained by fixed and random effects) R2. For each species, we fitted two candidate models – one that included a linear term, and second that included both linear and quadratic term of proportion of failure years as predictors. The best one was chosen based on the standard AIC criteria and only that one is reported.

Response: synchrony (S)

Species	Predictor: proportion of zeros	β	SE	<i>z</i> -statistic	<i>p</i> -value	R2m	R2c
<i>P. pinea</i>	linear	-0.007	0.001	-4.66	< 0.001	0.06	0.99
<i>A. scaphoides</i>	linear	-0.01	0.003	-3.58	0.001	0.12	0.80
<i>S. aucuparia</i>	linear	-0.01	0.001	-10.34	< 0.001	0.50*	
<i>Q. ilex</i>	linear	-2.11	0.55	-3.86	< 0.001	0.12	0.96
	quadratic	-1.39	0.54	-2.58	0.01		
<i>Q. humilis</i>	linear	-0.003	0.002	-1.47	0.142	0.01	0.95
<i>Q. rubra</i>	linear	-1.58	0.47	-3.36	< 0.001	0.50	0.94
		-0.94	0.43	-2.17	0.03		
<i>Q. alba</i>	linear	-0.02	0.003	-4.39	< 0.001	0.61	0.93
<i>Q. montana</i>	linear	-0.009	0.004	-2.36	0.018	0.75 [#]	
<i>C. pallens</i>	linear	-4.23	0.49	-8.68	< 0.001	0.97 [#]	
	quadratic	-2.63	0.42	-6.26	< 0.001		
<i>C. macra</i>	linear	-2.71	0.46	-5.86	< 0.001	0.95 [#]	
	quadratic	-1.96	0.42	-4.70	< 0.001		
<i>C. lyallii</i>	linear	0.04	0.01	3.43	< 0.001	0.13	0.99
<i>P. tenax</i>	linear	-0.58	0.22	-2.61	0.009	0.61*	
	quadratic	-0.93	0.24	-3.92	< 0.001		

* Plants were monitored at one site, thus a model without site as random effect was fitted for that species. The R-squared was calculated as Pearson's correlation coefficient between the observed and predicted values of the response.

[#] Conditional R2 could not be computed due to low variance of the random effect.

Table S5. Summary of linear mixed models regressing the variation in the plant CVi vs that plant fecundity (**prediction 5**). R2m is the marginal (i.e. the proportion of variance explained by fixed effects) and R2c is the conditional (i.e. the proportion of variance explained by fixed and random effects) R2. For each species, we fitted three candidate models – one that included a linear term of tree-level mean fecundity as a predictor, a second that included both linear and quadratic term, and the third one that fitted the relationship between the response and predictor as a Self-Starting asymptotic function. The best one was chosen based on the standard AIC criteria and only that one is reported.

Response: CVi

Species	Predictor: fecundity	β	SE	<i>z</i>-statistic	<i>p</i>-value	R2m	R2c
<i>P. pinea</i>	asymptotic	1.04	0.10	10.20	< 0.001	0.41	0.50
<i>A. scaphoides</i>	asymptotic	0.65	0.06	10.06	< 0.001	0.20	0.38
<i>S. aucuparia</i>	asymptotic	1.0	0.09	11.46	< 0.001	0.56*	
<i>Q. ilex</i>	asymptotic	1.02	0.12	8.45	< 0.001	0.25	0.26
<i>Q. humilis</i>	asymptotic	1.01	0.08	12.05	< 0.001	0.44	0.51
<i>Q. rubra</i>	asymptotic	0.90	0.14	6.33	< 0.001	0.49	0.57
<i>Q. alba</i>	asymptotic	1.01	0.24	4.14	< 0.001	0.25	0.31
<i>Q. montana</i>	asymptotic	0.99	0.15	6.44	< 0.001	0.57 [#]	
<i>C. pallens</i>	asymptotic	0.96	0.09	10.90	< 0.001	0.36	0.41
<i>C. macra</i>	asymptotic	1.07	0.10	10.37	< 0.001	0.51	0.52
<i>C. lyallii</i>	asymptotic	2.67	0.57	4.71	< 0.001	0.21	0.27
<i>P. tenax</i>	asymptotic	21.90	3.87	5.66	< 0.001	0.68*	

* Plants were monitored at one site, thus a model without site as random effect was fitted for that species. The R-squared was calculated as Pearson's correlation coefficient between the observed and predicted values of the response.

[#] Conditional R2 could not be computed due to low variance of the random effect.

Table S6. Summary of generalized mixed models regressing the variation in the plant reproductive synchrony (S) vs that plant fecundity (**prediction 6**). R2m is the marginal (i.e. the proportion of variance explained by fixed effects) and R2c is the conditional (i.e. the proportion of variance explained by fixed and random effects) R2. For each species, we fitted two candidate models – one that included a linear term, and second that included both linear and quadratic term of tree-level mean fecundity as a predictor. The best one was chosen based on the standard AIC criteria and only that one is reported.

Response: synchrony (S)

Species	Predictor: fecundity	β	SE	<i>z</i> -statistic	<i>p</i> -value	R2m	R2c
<i>P. pinea</i>	linear	0.02	0.005	4.26	< 0.001	0.11	0.99
<i>A. scaphoides</i>	linear	5.33	0.73	7.34	< 0.001	0.40	0.90
	quadratic	-1.69	0.67	-2.52	0.012		
<i>S. aucuparia</i>	linear	5.22	0.56	9.39	< 0.001	0.63*	
	quadratic	-1.91	0.50	-3.85	< 0.001		
<i>Q. ilex</i>	linear	0.9e-03	0.3e-03	3.47	< 0.001	0.22	0.96
<i>Q. humilis</i>	linear	0.001	0.0009	1.58	0.114	0.01	0.95
<i>Q. rubra</i>	linear	0.10	0.02	4.35	< 0.001	0.56	0.96
<i>Q. alba</i>	linear	0.15	0.03	5.74	< 0.001	0.70	0.96
<i>Q. montana</i>	linear	0.08	0.05	1.80	0.072	0.76	0.80
<i>C. pallens</i>	linear	0.08	0.009	8.42	< 0.001	0.97	0.98
<i>C. macra</i>	linear	0.07	0.02	4.64	< 0.001	0.78	0.96
<i>C. lyallii</i>	linear	0.31	1.46	0.21	0.834	0.17	0.99
	quadratic	3.56	1.31	2.26	0.01		
<i>P. tenax</i>	linear	0.08	0.02	4.84	< 0.001	0.62*	

* Plants were monitored at one site, thus a model without site as random effect was fitted for that species. The R-squared was calculated as Pearson's correlation coefficient between the observed and predicted values of the response.

Conditional R2 could not be computed due to low variance of the random effect.

Table S7. Summary of generalized mixed models regressing the variation in pre-dispersal seed predation vs log-transformed mean plant fecundity (**prediction 7**). R2m is the marginal (i.e. the proportion of variance explained by fixed effects) and R2c is the conditional (i.e. the proportion of variance explained by fixed and random effects) R2. For each species, we fitted four candidate models, including mean plant fecundity as a linear or quadratic term, with AR1 temporal correlation structure either included or excluded. The best model was chosen based on the standard AIC criteria and only that one is reported.

Response: proportion of predated seeds

Species	Predictor: fecundity	β	SE	AR1	<i>z</i> -statistic	<i>p</i> -value	R2m	R2c
<i>P. pinea</i>	linear	-0.16	0.11	0.37	1.40	0.163	0.01	
<i>S. aucuparia</i>	linear	0.12	0.20	0.13	0.62	0.533	0.01	
<i>Q. ilex</i>	linear	1.02	0.17	0.03	5.99	< 0.001	0.42	
<i>Q. humilis</i>	linear	1.67	0.26	0.08	6.35	< 0.001	0.60	0.63
<i>Q. rubra</i>	linear	-0.24	0.19	0.11	-1.25	0.211	0.01	
<i>Q. alba</i>	linear	0.28	0.23	-0.17	1.24	0.213	0.01	
<i>Q. montana</i>	linear	-0.04	0.29	-0.38	-0.15	0.878	0.00	

Table S8. Annual variability and synchrony of seed production in studied species. Mean CVi is the average individual-plant coefficient of variation. S is the mean synchrony of seed production by plants, measured by the average pairwise correlation of seed production of individual plants in a population through time. %fruit90 is the proportion of all fruits produced by the top 10% most fecund individuals. N plant is the number of plants monitored, N years is the number of years a population was monitored.

Species	N plant	N year	Below 90 th percentile of total seed production		Super-producers: Plants above 90 th percentile of total seed production			
			CVi	S	N plants	CVi	S	%fruit90
<i>P. pinea</i>	187	13	1.25	0.70	19	0.77	0.80	41%
<i>A. scaphoides</i>	507	28	3.08	0.39	51	2.08	0.56	40%
<i>S. aucuparia</i>	299	20	1.91	0.47	30	1.64	0.76	42%
<i>Q. ilex</i>	225	12	2.39	0.54	23	2.01	0.80	48%
<i>Q. humilis</i>	172	12	2.59	0.33	18	1.94	0.46	53%
<i>Q. rubra</i>	44	16	2.01	0.46	5	1.64	0.84	25%
<i>Q. alba</i>	51	16	2.38	0.36	5	1.83	0.69	28%
<i>Q. montana</i>	33	18	2.20	0.39	4	1.96	0.47	39%
<i>C. pallens</i>	217	30	2.69	0.69	22	2.04	0.88	30%
<i>C. macra</i>	125	28	2.79	0.66	13	2.11	0.84	32%
<i>C. lyallii</i>	94	24	3.82	0.90	10	3.26	0.94	40%
<i>P. tenax</i>	37	18	0.93	0.79	4	0.88	0.87	14%