

Conflicting selection from fire and seed predation drives fine-scaled phenotypic variation in a widespread North American conifer

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Recent work has demonstrated that evolutionary processes shape ecological dynamics on relatively short timescales (eco-evolutionary dynamics), but demonstrating these effects at large spatial scales in natural landscapes has proven difficult. We used empirical studies and modeling to investigate how selective pressures from fire and predispersal seed predation affect the evolution of serotiny, an ecologically important trait. Serotiny is a highly heritable key reproductive trait in Rocky Mountain lodgepole pine (*Pinus contorta* subsp. *latifolia*), a conifer that dominates millions of hectares in western North America. In these forests, the frequency of serotiny determines postfire seedling density with corresponding community- and ecosystem-level effects. We found that serotinous individuals have a selective advantage at high fire frequencies and low predation pressure; however, very high seed predation shifted the selective advantage to nonserotinous individuals even at high fire frequencies. Simulation modeling suggests that spatial variation in the frequency of serotiny results from heterogeneity in these two selective agents. These results, combined with previous findings showing a negative association between the density of seed predators and the frequency of serotiny at both landscape and continental scales, demonstrate that contemporary patterns in serotiny reflect an evolutionary response to conflicting selection pressures from fire and seed predation. Thus, we show that variation in the frequency of a heritable polygenic trait depends on spatial variation in two dominant selective agents, and, importantly, the effects of the local trait variation propagate with profound consequences to the structure and function of communities and ecosystems across a large landscape.

Greater Yellowstone Ecosystem | geographic selection mosaics | genes to ecosystems

Recent work has attempted to unify processes acting across spatiotemporal scales by connecting smaller-scale (e.g., individuals, patches) ecological processes to large-scale (e.g., ecosystems, landscapes) patterns via evolutionary mechanisms (1, 2). When heritable traits in foundation species have important community and ecosystem effects, ecological drivers that exert natural selection on these traits can propagate to large-scale patterns in community and ecosystem function, resulting in landscape heterogeneity driven by small-scale ecological variation (1, 3). Although there is a clear conceptual link between selection acting on individuals and landscape-scale patterns, making mechanistic connections across these very different scales has proven challenging (refs. 4 and 5; but see ref. 6). Forested ecosystems in particular are difficult to study, because long generation times prevent direct observations of any response to selection. Nevertheless, attempts to elucidate the connections between individual-level genetic variation and ecosystem and landscape structure are imperative, especially given the unprecedented pace of global change.

Here, we integrate empirical studies and simulation modeling to show that landscape structure in Rocky Mountain lodgepole pine (*Pinus contorta* subsp. *latifolia*) forests results from the combination

of opposing selective pressures from fire and a seed predator. We present empirical data relating reproductive success of individual trees to variation in selection exerted by a seed predator, as well as both empirical data and process-based simulations linking local selective pressure from fire and seed predation to landscape-scale variability in phenotype frequencies. We focus on serotiny, which is the retention of seeds in a canopy seed bank until high temperatures (typically stand-replacing fire) triggers seed release (7). Serotiny is an adaptive trait that increases regeneration following stand-replacing fires, and it is common among woody plants in fire-prone ecosystems (7). Lodgepole pine dominates millions of hectares in western North America, and serotiny in lodgepole pine is highly heritable (8) and has far-reaching ecological effects due to a strong correlation between prefire frequency of serotiny and postfire seedling density at local (i.e., patch) scales (9–11). The community- and ecosystem-level effects of variable seedling density are considerable in this system and include competitive exclusion of other trees (12), succession (10), and annual net primary production (13).

Most studies of the evolution of serotiny have focused on the relationship between fire regime and serotiny (7, 14), and both empirical and theoretical work suggest that serotiny is most common where stand-replacing fires occur predictably within the lifetime of the species (14–16). It follows that much geographical variation in serotiny is due to variation in fire regimes (17–19). Postdispersal seed predation may be another mechanism favoring the evolution of serotiny if delayed and synchronous seed release limits the abundance of postdispersal predators (7, 20). However, escape from postdispersal predation may come at the expense of extreme losses to predispersal predators that can easily access the canopy seed bank (21, 22), potentially exerting selection favoring increased allocation to seed defense (21, 23–25).

Significance

We investigate selective agents acting on serotiny, a polygenic trait with high heritability and well-documented effects on community and ecosystem structure and function. We show that the frequency of serotiny in lodgepole pine across Yellowstone National Park, and likely over large portions of the Rocky Mountains, represents the balance of selection by two agents: fire and a seed predator, the American red squirrel. Thus, ecosystem structure and function in these ecosystems likely varies as a result of spatial variation in these two selective agents.

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Previously, we demonstrated that the frequency of serotiny in lodgepole pine in the Greater Yellowstone Ecosystem (GYE) is negatively correlated with the abundance of a predispersal seed predator, the American red squirrel *Tamiasciurus hudsonicus* (26). In lodgepole pine forests, red squirrels are territorial central-place seed predators (27). Squirrels rely on large caches of lodgepole pine cones to survive the winter, and because entire cones are removed and stored intact, seed dispersal as a result of squirrel foraging is negligible (28). Previous work has shown that energy intake rates on nonserotinous cones are slightly higher than on serotinous cones, implying that preferential feeding on nonserotinous cones is a more profitable strategy (21). However, red squirrels cache nonserotinous cones almost exclusively during several weeks in the fall between seed maturation and when cones open and seeds are dispersed. Serotinous cones, however, are available year-round. This increased availability may result in greater numbers of serotinous cones being removed by squirrels and driving selection against serotiny. However, we lacked data linking this mechanism to the correlations between the frequency of serotiny and squirrel density observed in our field studies (26).

In the present study, we expand our previous work to investigate how hypothesized evolutionary mechanisms (stand-replacing fire and predispersal seed predation) drive landscape-scale variation in the frequency of serotiny in lodgepole pine in the GYE. We used natural variation in the abundance of seed predators to test the hypothesis that predispersal seed predation exerts selection against serotiny and to quantify predation rates. We then developed a stand-level process-based model to explore how the fitness of serotinous and nonserotinous trees varied in relation to variation in both fire frequency and seed predation. The use of simulation modeling allowed us to address evolutionary questions despite the long life span of lodgepole pine, and allowed us to directly compare the response to selection from processes operating on vastly different spatial and temporal scales.

Results

Selection from Seed Predation. We measured cone removal rates as an assay of seed losses due to predispersal predation for 216 serotinous and 207 nonserotinous trees at five sites. To address local (i.e., per-tree) squirrel density, we used the distance to the nearest squirrel territory center (identifiable by the presence of a large midden of chewed and cached cones) as a proxy for density. Because red squirrels are strongly territorial central-place foragers (27), the average distance to a midden decreases with increasing squirrel density (see *Materials and Methods* for additional details). We then used a generalized linear mixed model (GLMM) to evaluate cone survival in relation to cone type (first-year serotinous, >1-y-old serotinous, or nonserotinous) and distance to the nearest midden. To avoid confounding selection for cone type with selection based on tree size or age, we included diameter at breast height (DBH) as a covariate. Mean survival was lowest for first-year serotinous cones (0.81), followed by nonserotinous (0.87) and >1-y-old serotinous (0.97) cones. Survival increased with increasing distance from the nearest midden, and the slopes varied with cone type (Fig. 1). The slope of the relationship for serotinous cones was significantly greater than the slope for nonserotinous cones ($P < 0.05$), indicating that the difference in predation rates between serotinous and nonserotinous cones increased with increasing squirrel density (see *Table S1* for full model formulation and parameter estimates). Furthermore, at <50 m from the midden [where most seed predation occurs (27)], when accounting for all predation (i.e., from both first-year and older cones; see *Materials and Methods*), serotinous trees consistently experienced greater losses from predation than nonserotinous trees, and the difference increased with increasing squirrel density (Fig. 1).

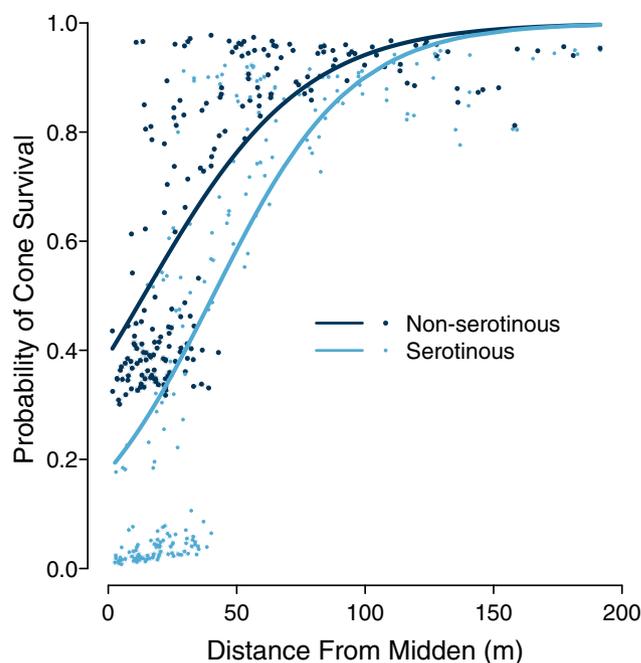


Fig. 1. Cone survival decreased more rapidly for serotinous ($n = 216$) than for nonserotinous ($n = 207$) trees with decreasing distance from the nearest midden (i.e., increasing squirrel density). Survival probabilities for serotinous cones include the effects of multiple years of predation. The fitted lines are predictions from a binomial generalized linear model.

Simulation Results. We developed a simulation model, based on a logistic population growth model, that integrated selection from both stand-replacing fire and seed predation by squirrels. Fires were implemented as either deterministic, recurring events or as a stochastic process, with return intervals varying within a realistic range for the GYE [100–400 y (29)]. Squirrel densities ranged from 0 to 1.5 individuals per ha (26) and were assumed to be constant within individual simulation runs (see *Discussion* for information about this assumption) (28, 30–32). In deterministic simulations, serotinous individuals were favored when squirrels were absent and the fire return interval was <350 y. Increasing selection from seed predation decreased the relative fitness of the serotinous genotype (defined here as the mean population fitness of serotinous individuals/the mean population fitness of nonserotinous individuals) and resulted in higher fire frequencies being required before serotiny was favored (i.e., relative fitness, ≥ 1 ; Fig. 2).

In stochastic simulations with no seed predation, the frequency of serotiny declined monotonically with increasing fire return interval (Fig. 3A). We observed similar patterns after incorporating seed predation, and the predicted frequency of serotiny at a given fire return interval decreased with increasing squirrel density (Fig. 3B). At squirrel densities ≥ 1.5 individuals per ha, the model predicted no serotiny regardless of the fire regime. Model predictions were nearly identical to landscape-scale patterns we previously reported (26); we observed a steep negative correlation between squirrel density and serotiny at high fire frequencies, and little serotiny regardless of squirrel density at low fire frequencies (Fig. 4).

Discussion

Our results demonstrate a clear connection between two selective agents (fire and seed predators) and both the relative fitnesses of serotinous and nonserotinous individuals (Figs. 1 and 2) and their frequencies at the stand level (Fig. 4) in lodgepole pine. Combined with previous work showing that serotiny is

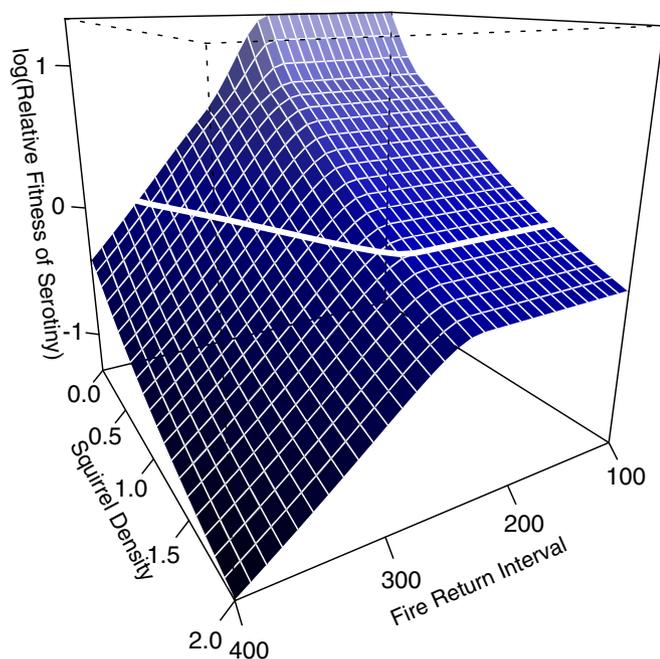


Fig. 2. Simulation results showing how the relative fitness of the serotinous genotype varied with fire return interval (years) and squirrel density (individuals per hectare). Serotiny was favored (above the white line; log relative fitness, >0) with short fire return intervals and low squirrel densities, whereas nonserotinous individuals were increasingly favored even at high fire frequencies as squirrel density increased.

polygenic and highly heritable (8), that stand-level serotiny covaries with both fire frequency and squirrel density at landscape scales (26), and that serotiny has significant effects on community and ecosystem structure and function (9–11), we argue that the effects of selection on serotiny propagate across multiple scales, driven by forcing mechanisms that are operating at relatively large (i.e., hundreds of hectares for fires) and small (~ 1 –20 ha for squirrels) scales to influence landscape structure (Fig. S1). Such cross-scale dynamics are well supported by theory but are difficult to demonstrate empirically (5, 33, 34). It is unclear how common such dynamics are in natural systems; however, our results suggest that large-scale “noise” in ecological systems (e.g., variation in the frequency of serotiny) may be attributable to complex interactions at multiple spatial scales, and that evolutionary change in response to local variation in selective agents can be an important driver of these spatial patterns.

Much recent work has focused on investigating how evolutionary processes shape ecological dynamics (1, 5, 35), and on how resulting ecosystem-level dynamics may produce feedbacks and affect the selective environment of foundation species (36). However, demonstrating the complete chain of eco-evolutionary relationships in natural ecosystems has proven challenging (5, 37). In lodgepole pine, serotiny has important ramifications from individual to ecosystem levels (Fig. S1). At the individual level, serotiny determines when seeds are dispersed and the exposure to predispersal seed predators, and thus is tightly associated with fitness. Serotiny is also highly heritable; although exact estimates of heritability are unavailable for this species, a recent association mapping study demonstrated that 11 genomic regions accounted for $\sim 50\%$ of the variation in serotiny in lodgepole pine (8).

The combination of high heritability and tight coupling to fitness suggests that serotiny is likely to be a target of selection and to respond rapidly to changes in selection with resulting shifts in the frequency of serotiny in the population. At the community and ecosystem levels, the degree of prefire serotiny affects postfire

seedling density. Following widespread fires in the GYE, lodgepole pine sapling density varied over several orders of magnitude (0–500,000+ seedlings per ha), and this variation was most strongly explained by the prefire frequency of serotiny (9–11, 13). Unsurprisingly, this extreme variation in sapling density had significant community- and ecosystem-level consequences as well (9, 11–13). Sparse recovery may allow for the colonization of other ecosystem types that would otherwise be outcompeted, increasing landscape heterogeneity (12). Recovery density may also affect successional

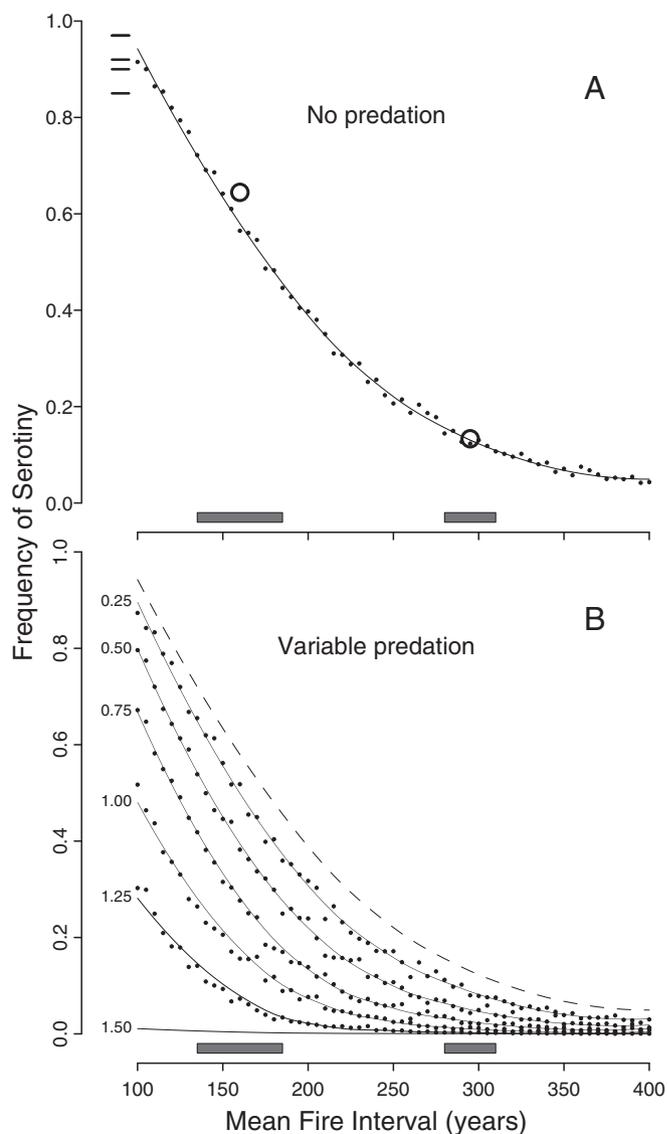


Fig. 3. Simulations predicted a decline in the frequency of serotiny with increasing fire return intervals in the absence of squirrels (A), and a decrease in the frequency of serotiny with increasing squirrel density (B). Points show the mean of 1,000 model runs at a given parameter combination, and the fitted lines are predictions from LOESS models. The gray bars near the x axis show the ranges of fire intervals for low (135–180 y) and high (280–310 y) elevations in the GYE (29). In A, the open circles are the frequencies of serotiny at these elevations in the absence of squirrels predicted from a regression on empirical data (Fig. 4; ref. 26), showing very close correspondence between empirical results and simulation predictions. The small horizontal lines in the *Upper Left* of A show frequencies of serotiny in isolated mountain ranges that lack squirrels (39). Fire return intervals for these ranges are likely <150 y. The solid lines in B show predictions at different squirrel densities (indicated by small numbers at the left side of the plot); the dashed line shows predictions in the absence of squirrels.

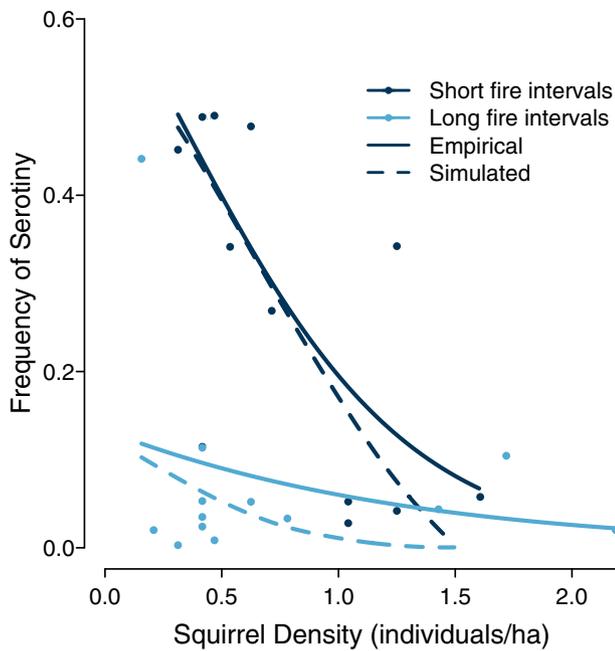


Fig. 4. Simulation predictions (dashed lines) were qualitatively very similar to regression predictions for empirical data (points and solid lines) from ref. 26. At sites with short mean fire intervals (135–185 y; dark colors), serotiny is variable, and both simulated and empirical data show a negative correlation between the frequency of serotiny and squirrel density. At long fire intervals (280–310 y; light colors), serotiny is uncommon and the relationship between squirrels and the frequency of serotiny is weak. The figure is modified from ref. 26 and used with permission.

processes (10), and postfire annual net primary productivity and total leaf area are strongly associated with lodgepole pine sapling density (13). Thus, in lodgepole pine forests, obtaining a clear understanding of ecosystem dynamics at scales ranging from individuals to landscapes requires analysis of the evolutionary mechanisms driving variation in serotiny.

Our model allows for the manipulation of fire frequency and observation of subsequent stand-level evolutionary responses over long timescales; observations of these dynamics would be difficult or impossible in the field. Given the well-known effects of fire on the evolution of serotiny (7, 14–17), such observations are required to understand the evolutionary effects of seed predation in context, and to demonstrate that selection observed in the field (Fig. 1) results in evolution (Figs. 2 and 3) and drives spatial patterns in serotiny [Fig. 4 (26)]. The model makes the simplifying assumption that squirrel densities are constant in time for a given scenario. Although this is certainly unrealistic, the available data demonstrate that squirrel populations are stable over short to medium timescales (i.e., years to decades). In part, this is driven by stable cone production of lodgepole pine, which is the primary food source for red squirrels in this system (21, 30–32). The strictly territorial habit of red squirrels also promotes population stability, as vacant territories are rapidly reoccupied and overwinter survival of squirrels lacking territories is extremely low (28, 31). Therefore, fluctuations in density are likely to be small; adding such fluctuations to the model would increase the variance, but have little impact on the overall conclusions. However, we lack information on fluctuations over longer timescales (potentially in response to changes in postfire recovery as the stand evolves), and the density of squirrel territories may be more variable in ecosystems with less stable cone production (28). If changes in the frequency of serotiny over time lead to changes to stand structure that ultimately affect squirrel population densities, then there is the potential for feedbacks in the system that could either reinforce or dampen spatial

patterns in selection on serotiny. Alternatively, stand structure may be more influenced by factors such as bedrock (38), which could lead to similar stand structure from one tree generation to the next, resulting in consistent spatial variation in both squirrel densities and selection.

Evolutionary Mechanisms. Previous studies have focused on the role of fire in driving the evolution of serotiny (7, 17–19), although few have addressed seed predators as alternative selective agents (but see refs. 7, 19, and 39), and ours is the first (to our knowledge) to demonstrate strong selection from seed predation driving local-scale (i.e., stand-level) variation in serotiny. In opposition to these processes are potentially high rates of gene flow in lodgepole pine that may have a homogenizing effect on spatial variation in serotiny (40); however, strong selection can drive local adaptation despite the presence of gene flow (41–45), and even at small spatial scales, local adaptation is possible in the presence of gene flow (46). It is likely that gene flow could explain some of the variation in serotiny in this system. However, our simulations predicted strong selection and evolutionary responses to both fire and seed predation in the absence of gene flow, resulting in less serotiny when fires were uncommon or seed predation was more intense (Figs. 2 and 3). Furthermore, the predictions of the model were very similar to landscape-scale patterns we observed in the GYE (Fig. 4), suggesting that selection is the dominant force affecting the local frequency of serotiny in this system. It is likely that a spatially explicit version of the model that directly simulated gene flow would demonstrate weaker evolutionary responses overall, with more stable coexistence of the two types due to the continuous dispersal of maladapted individuals into a given region. In contrast, our spatially implicit model showed a tendency toward fixation of either the serotinous or nonserotinous type when selection was strong, a situation that is rarely observed in the field (9, 26).

Muir and Lotan (47) found that the frequency of serotiny in lodgepole pine was most strongly related to the type of disturbance that initiated a stand, with fire-initiated stands being much more serotinous than other stands. This suggests the potential for very rapid change in the frequency of serotiny (i.e., a single fire interval), and that high frequencies of stand-replacing fire should result in high frequencies of serotiny (because most stands will have initiated due to a stand-replacing fire). Our simulation results generally support this interpretation. In very old stands with dynamics dominated by density-independent mortality, the frequency of serotiny was reduced compared with postfire stands. However, our simulations also suggest that the legacy of long fire return intervals persists even if stands were initiated by fire; i.e., fire-initiated stands that experienced short mean fire return intervals had higher serotiny than fire-initiated stands that experienced long fire return intervals (Fig. 3). Furthermore, the effects of seed predation were completely dominant at very high squirrel densities (>1.5 individuals per ha). Many trees show local adaptation and rapid (per-generation) responses to selection (45, 48); our results in lodgepole pine are consistent with these observations. Lodgepole pine forests have probably been present in the GYE for ~10,000 y (49). Our simulations, which were run for 2,000 y, suggest that this is sufficient time for consistent and sometimes strong divergent selection to cause local adaptation and spatial structure in serotiny across the landscape.

Implications for Global Change. The rapid pace of global change is likely to have widespread evolutionary effects due to changing strength and direction of biotic and abiotic interactions (50). In addition to relatively predictable dynamics (e.g., selection favoring greater dispersal or greater tolerance to novel local conditions), these changes may be complex and nonlinear, in part due to interactions between ecological and evolutionary processes (2, 51). For example, in lodgepole pine forests, increasing fire frequency due to forecasted increases in the incidence of fire weather (52)

may result in serotiny being increasingly favored regardless of squirrel density, or may result in complex changes due to non-linear responses to repeated burning (53). Mechanistic models incorporating eco-evolutionary dynamics will be increasingly important in predicting these responses, especially when global change results in combinations of selective agents (e.g., fire frequency and squirrel density in our system) that are outside the scope of current knowledge.

Conclusions

Our results demonstrate that squirrels select against serotiny and that the strength of selection increases with increasing squirrel density, supporting previous results showing negative associations between red squirrel density and the frequency of serotiny in lodgepole pine at landscape and regional scales (26, 39). We have also shown that a simple population model incorporating selection from fire and seed predation reproduces empirical observations (Figs. 3 and 4), suggesting that stand-level frequencies of serotiny in lodgepole pine represent the outcome of selection from fire (favoring serotinous trees) and seed predation (favoring nonserotinous trees). Although gene flow is extensive in lodgepole pine (40), strong selection can overwhelm gene flow and cause local divergence (41–45). In lodgepole pine forests in the GYE, we found little evidence that gene flow prevents local adaptation; rather, our results suggest that divergent selection is the dominant factor affecting spatial variation in the frequency of serotiny. These findings have implications for ecosystem processes due to the influence of serotiny on stand-level responses to fire (9, 10), as well as for global change in a variety of ecosystems, because changes in selective regimes may produce unexpected effects to ecosystem structure and function.

Materials and Methods

Study Area. We performed all field work in mature (i.e., not burned within 100 y) lodgepole pine forests within the GYE, Wyoming, United States. We selected five sites that were known to vary from ~0.5 to 1.5 squirrels per ha (26). The sites ranged from 25.6 to 45.4 ha and were dominated (>90%) by lodgepole pine. One site was sampled in July 2010 and July 2011, and four sites were sampled in August 2011 and July to August 2012.

Cone Survival. We estimated survival probabilities of serotinous and nonserotinous cones by repeated observation with a one-year (± 1 mo) intervening interval. During the initial visit, we photographed the tops of randomly selected, individually marked trees ($n = 216$ serotinous and 207 nonserotinous trees). We also measured the DBH of each tree. During subsequent visits, we recreated the initial photograph, using the same position, camera angle, and focal length. We compared these photographs in the laboratory and tabulated the number of cones that had disappeared during the 1-y interval [i.e., individual cones that were present in the first photograph but absent in the second (Dataset S1)]. A cone was counted as surviving if it could be clearly identified (based on position relative to other cones, branch features, etc.) in both photographs. Cones were counted as absent if the cone's location in the second photograph could be clearly seen (i.e., was identifiable based on nearby cones and branches and not obscured). Cones that were not identifiable in this manner were treated as missing data and ignored for analysis. The rate of disappearance between intervals is an appropriate measure of the predation rate from red squirrels because no other organism removes cones from lodgepole pine (due to extreme defense requiring specialized predators), and cones do not generally fall from branches (even after they have opened) (21, 25). Because survival of serotinous cones is likely to vary as the cone ages, we analyzed survival as a function of the degree of weathering,

with separate estimates for green (first-year), brown (likely 1- to 5-y-old), and gray (>5-y-old) cones. Exploratory analysis revealed that survival of gray cones was nearly 100% in all circumstances, so we omitted these cones from our analyses.

To address how squirrel density affected cone survival, we used the distance of each tree to the center of the nearest squirrel territory as a proxy for density. Territory centers were located by an exhaustive ground search by two observers and were identified by the presence of a large, active midden containing burrows and discarded cone remnants (21). Because red squirrels are strictly territorial, the average distance to a midden decreases with increasing density; for four sites where mean squirrel density for the entire site was known (from ref. 26), squirrel density and the distance from trees to the nearest midden were strongly negatively correlated ($r = -0.88$).

We used package lme4 in R to develop GLMMs to analyze survival probability as a function of distance to the midden (54). All models used the number of surviving cones and the number of cones initially observed as the response variable and assumed binomial errors and a logit link, and all included site as a random variable. Fixed effects consisted of cone type, distance to the midden, and an interaction between the two. For the relationship between distance and survival, we developed three models: fixed slopes and random intercepts, random slopes and intercepts, and no effect of distance. We also tested models including or excluding DBH as a covariate, for a total of six models. We compared all models using the Bayesian information criterion (BIC). The fully parameterized model minimized the BIC. Furthermore, $\Delta\text{BIC} \geq 39$ for all other models, indicating little uncertainty in which was the best model (55); therefore, we selected the full model alone for interpretation (see Table S1 for parameter estimates). Because serotinous cones are exposed to predation for multiple years, the overall cone survival is the product of survival probabilities over the years in which cones are exposed to predation. Thus, in addition to the GLMM described above, we calculated for each serotinous tree the probability that a cone survived long enough to enter the "weathered" category (at which point survival was nearly 100%), and then compared this to the nonserotinous survival rate as a function of distance using a logistic regression (Fig. 1).

Simulation Model. We developed a stand-level simulation model to provide an integrated estimate of the relative fitness of both tree types and to address the potential for the observed predation rates to drive variation in the frequency of serotiny. Our general approach was to use a density-dependent population model (56) to simulate stand development, starting with postfire initiation and proceeding through self-thinning (or infilling, for sparse stands) and stand maturity. The response of lodgepole pine to fire is well known; therefore, we used the literature to parameterize this portion of the model. Fires were initiated at regular intervals (for deterministic scenarios) or probabilistically (for stochastic scenarios), and we varied either the fire return interval or the fire probability to assess the effect of fire frequency on serotiny in the simulated stand. For predation, we used the results presented in this paper to reduce the reproductive success of trees as a function of serotinous/nonserotinous state, and varied the intensity of predation based on the ranges observed in the field. See *SI Text* and Table S2 for complete details on model specification and parameterization. Sensitivity analysis was conducted by examining how the relationship between serotiny and both fire and seed predation was affected by perturbations to key model parameters (*SI Text*, Figs. S2–S7).

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Supporting Information

Talluto and Benkman 10.1073/pnas.1400944111

SI Text

Simulation Description and Sensitivity Analysis

Model Description. The central structure of the model was a modification of the Ricker model, a discrete-time density-dependent population model (1). At each annual time step, the model tracked three state variables: the density of serotinous ($n_{s,t}$) and nonserotinous ($n_{n,t}$) individuals (and, by extension, the total population size: $N_t \equiv n_{s,t} + n_{n,t}$), and the current stand age (a_t). Serotinous individuals produced and accumulated seeds, with dispersal allowed only immediately following a fire, whereas nonserotinous individuals dispersed seeds during each time step. We assumed that both phenotypes had identical mortality and seed production rates. We modeled serotiny as a single-locus haploid trait. Although this is certainly an unrealistic assumption (2, 3), we chose this method for simplicity and computational efficiency. For deterministic scenarios, we analyzed the relative fitness of serotiny after a single generation of selection. For stochastic scenarios, we present the means of 1,000 model runs (of 2,000 model-years each) at each parameter combination. Because the frequency of serotiny varies with stand age (as nonserotinous trees fill in gaps and adult trees die), we analyzed the frequency of serotiny at a constant stand age of 100 y. A complete list of state variables and parameters, along with the results of parameterization, is available in Table S2.

Population Dynamics. Population dynamics varied depending on whether a stand was mature (i.e., stands where self-thinning had ceased; $a_t < a_{mature}$, the age at maturity, estimated at 200 y from ref. 4) or immature. For immature stands, we modified the Ricker model:

$$n_{s,t+1} = n_{s,t} e^{\rho \left(1 - \frac{N_t}{K(a_t)}\right)}; \quad 0 < a_t < a_{mature}, \quad [\text{S1}]$$

where ρ is the population growth rate and K is the self-thinning function, representing the carrying capacity for a given stand age. We used an identical corresponding equation for nonserotinous individuals ($N_{n,t+1}$). We lacked specific literature values for ρ . Therefore, we used preliminary model runs to tune this parameter, using infilling time as a relatively independent criterion for tuning. Literature estimates suggest that stands that are sparse at initiation reach high density in ~50 y (5, 6). We selected $\rho = 0.05$ as a value that replicated this behavior (stands achieved >60% of maximal density in 50 y). We discuss the implications of this tuning further in the section on sensitivity analysis. Finally, empirical data show that lodgepole pine continues to establish for several decades following fires (5, 6), and some of this establishment is likely due to seeds released from immature trees that are genetically serotinous but produce open cones at a young age (2, 7). Lotan and Perry (8) speculated that cone opening by immature serotinous trees is an adaptive trait that allows serotinous trees to contribute to ongoing postfire recovery in stands where initial recovery was sparse. Therefore, for stands younger than 30 y old, we assumed infill trees were serotinous or nonserotinous in proportion to the existing population. Once the stand aged beyond 30 y, we assumed serotinous trees produced closed cones and all infilling trees were from nonserotinous parents.

Carrying capacity was determined using a self-thinning curve where maximum stand density declined as a function of age using a power rule (9):

$$\log K(a) = \beta_0 + \beta_1 \log a, \quad [\text{S2}]$$

where β_0 and β_1 are parameters fit from empirical data. For self-thinning, density is generally predicted as a function of tree size; here, for simplicity we assumed that individual size increases with stand age, and modeled density directly as a function of stand age. We fit the parameters of this model directly using empirical data (10) ($n = 4$, stands ranging from 12 to 250 y old, $R^2 = 0.997$). Our empirically estimated slope of -1.39 is similar to the -1.5 slope of Yoda's law (9), suggesting that the assumption that size scales with age is plausible.

For mature stands, we assumed the population was stable (i.e., all dead individuals were immediately replaced), with an annual deterministic mortality rate μ . We assumed annual turnover of 0.5% based on mortality rates for conifers in the region (11). Because we lacked information on the variance in mortality rates, and mortality rates in *Pinus* may be as high as 2% (11), we varied this parameter from 0.5% to 2% during sensitivity analysis.

Fire and Postfire Recovery. We used both deterministic and stochastic processes to govern fire occurrence. In both cases, fire occurrence was controlled by a single parameter F , the fire return interval in years. We used deterministic simulations to examine the relative fitness of serotiny under varying fire and predation rates (e.g., Fig. 2 in main text). For these simulations, a fire was initiated every F years. We used stochastic simulations to examine how model predictions about the frequency of serotiny compared with empirical results (Figs. 3 and 4 in main text). For these simulations, fires were initiated randomly with an annual probability of $1/F$. For all simulations, fires killed all individuals, triggered the release of stored serotinous seeds, and initiated the postfire recovery phase of the model. Fire was allowed in all stands exceeding 100 y old (12). We varied F from 100 to 400 y in simulations, based on published estimates of 135–185 y for low elevations and 280–310 y for high elevations within the Greater Yellowstone Ecosystem (GYE) (13).

We modeled postfire seedling density as a function of the prefire size of the canopy seed bank, s_t , given immigration rate (γ_0) and seedling density (γ_1) parameters (13–16):

$$\begin{aligned} \log n_{n,t} &= \gamma_0, & a_t &= 0. \\ \log n_{s,t} &= \gamma_1 s_{t-1}, \end{aligned} \quad [\text{S3}]$$

Because we lacked data on necessary processes (e.g., seed production, dispersal, and germination rates), we treated s_t as a latent variable determined by population density ($n_{s,t}$, $n_{n,t}$) and the seed predation rate (see below). This allowed us to parameterize the relationship using available data.

Seed Predation. We modeled seed predation as a proportional reduction in the size of the canopy seed bank (for serotinous individuals) or a reduction in the rate of infilling (for nonserotinous individuals). Empirical data show that the size of the canopy seed bank (represented by the prefire frequency of serotiny) strongly predicts postfire stand density (16, 17). Our model assumes that rates of seed predation measured for individual trees translates to a stand-level reduction in the canopy seed bank. Thus:

$$s_t = (1 - \pi_s) s_{max} \frac{n_s}{K(a_t)}, \quad [\text{S4}]$$

where π_s is the predation rate of cones on serotinous trees, s_{max} is the maximal size of the accumulated seed bank (i.e., when the

population is saturated and nonpredation seed losses are balanced by seed production), and $n_{s,t}/K(a_t)$ is the ratio of the current serotinous population size to the maximal population size (thus normalizing the seed bank to the number of individuals available to produce seeds). Because we lacked the data to express s_t and s_{max} in terms of individual seeds, we instead abstracted away processes of germination and seed dispersal, and expressed these quantities in terms of seedlings. We were then able to parameterize this relationship using data relating postfire seedling density to the frequency of serotiny [i.e., $n_{s,t}/K(a_t)$ assuming the population is at carrying capacity]. We built a regression model using data from refs. 16 and 17 to parameterize Eq. S3 using the relationship in Eq. S4, assuming the intercept (i.e., the number of seedlings predicted when no trees were serotinous) represented immigration of nonserotinous seeds dispersed from outside the focal (burned) stand. Because the sites for the source data were chosen without regard to squirrel density or seed predation (16, 17), we set π_s to the median predation rate observed in the present study for the purposes of parameterization, recognizing that this results in unexplained variance (due to predation rates varying among sites) but that the mean predictions will be unbiased assuming sites were selected without bias in the original study. Finally, we used the data in the main text of this paper to relate the predation rates (π_s , π_n) to squirrel density, and then varied squirrel density systematically across a range of realistic values for the GYE (18) to determine the effects of predation on selection on serotiny.

Seed Dispersal and Recruitment. The number of seeds dispersed and the contribution to the new class of seedlings in each model year depended on stand age, frequency of serotiny, and whether a fire occurred. In nonfire years, all seeds produced and not eaten by squirrels either entered the canopy seed bank or were immediately dispersed. For young stands (see *Population Dynamics* above), both serotinous and nonserotinous seeds were dispersed, whereas in all other stands all serotinous seeds entered the canopy seed bank. In fire years, seeds were dispersed as in nonfire years, but in addition all seeds in the canopy seed bank were also dispersed. Finally, dispersed seeds were converted into new individuals based on the number of new individuals predicted by the population model, with serotinous and nonserotinous individuals recruiting in proportion to their representation in the dispersed seed cohort. All extra seeds not recruited were lost (i.e., there was no soil seed bank).

Model Evaluation. We first ran the model with deterministic parameters to directly assess how variation in fire frequency and squirrel density affected the fitness of serotinous trees. To do this, we calculated the per-generation population growth rate for both genotypes (λ_s and λ_n for serotinous and nonserotinous trees, respectively), and then calculated the relative fitness of serotinous trees as λ_s/λ_n . To assess how these fitness differences might translate to observed frequencies of serotiny in real stands, where average fire regimes are known but exact fire intervals are variable, we used stochastic disturbances and varied fire probability and squirrel density. We ran 1,000 simulations at each parameter combination, and report the observed frequency of serotiny in 100-y-old stands (chosen to provide a fixed point for comparison of all model runs) after 2,000 y of simulation. For validation, we compared the model predictions with empirical results from our previous work (18).

Sensitivity Analysis. Model sensitivity was assessed by altering parameters one at a time and inspecting the effect of these perturbations on the relationship between the predicted frequency of serotiny and both fire frequency and the seed predation rate for stochastic scenarios. We evaluated sensitivity under two conditions: variable fire probability (ranging from 0.0025 to 0.01

and corresponding to expected intervals of 100–400 y, as in the main model runs) with no seed predation (i.e., squirrel density = 0), and variable squirrel density (range: 0–2.1 individuals per ha) with fire probability held constant at 0.00625 [i.e., expected 160-y intervals, approximating fire frequency at low elevations in the GYE (13)]. Under each of these conditions, we altered parameter values one at a time. For parameters lacking SE estimates (population growth rate and mortality rate), we chose four to five values from the ranges given in Table S2. For all other parameters, we chose values ranging from –2 to +2 SEs relative to the mean in increments of one-half the SE. Each combination was evaluated over 1,000 separate model runs as in the main analysis.

SI Results

Population Growth Rate. The model was sensitive to the population growth rate r , but only when the growth rate was very low (Fig. S2). When the growth rate was 0.01, serotiny was much more strongly favored than under the default, even at low fire frequency. Further exploration of low values of ρ ($0.01 \leq \rho \leq 0.1$) revealed a threshold of approximately $\rho = 0.03$; below this value, serotiny is favored at increasing rates. However, growth rates this slow also produced unrealistic estimates of infilling rates (e.g., >100 y to reach mature densities), suggesting that these values are not plausible (5, 6). At values higher than the default, population growth rate had little effect on the frequency of serotiny, with no change in the model predictions when $\rho \geq 0.11$.

Low growth rates favoring serotiny is unsurprising given the structure of our model. This parameter controls the rate at which nonserotinous seedlings fill in stands with sparse postfire recovery. Very low growth rates result in incomplete infilling, resulting in less competition for serotinous seedlings that established immediately after a fire and ultimately resulting in higher frequencies of serotiny in mature stands. The lack of sensitivity at high growth rates is likely the result of complete saturation of sparse stands with nonserotinous seedlings; beyond this saturation point, increasing the growth rate has no effect on postfire stand composition. However, our default value of $\rho = 0.05$ is close to a threshold, below which the model becomes highly sensitive to growth rate. If actual growth rates in lodgepole pine are lower than what we have used, then the predictions of our model may be poor. MacDonald and Cwynar (19) reported on doubling times in lodgepole pine populations undergoing postglacial expansion; the fastest-expanding population in their study had a population growth rate of 0.009, and the mean of all populations was 0.001, substantially lower than our default. However, our model simulates a burned stand in a connected forest, where competition is minimal, the climate is suitable, soil is mature (i.e., not postglacial), and seed sources from surviving trees and nearby stands are present. It is likely that growth rates under these conditions will be significantly greater than at the leading edge of population expansion that likely would be slowed by dispersal limitation, suboptimal climatic conditions, poorly developed soil, and competition with existing vegetation types.

Self-Thinning. The model showed little sensitivity to the self-thinning curve (β_0 and β_1). In the absence of seed predation, neither parameter had an effect on model predictions. When squirrel densities were varied, we observed some sensitivity to the self thinning curve at low to moderate squirrel densities (~ 0.7 – 1.5 individuals per ha for β_0 , 0.0 – 0.5 individuals per ha for β_1). At these densities, a large increase (2 SEs) in either the intercept (indicating higher densities at stand initiation) or the slope (indicating slower thinning over time) resulted in a decrease of as much as 30% in the predicted frequency of serotiny (Figs. S3 and S4).

Mortality Rate. Mortality rates above the rate of 0.005 used as our model default resulted in large decreases in the predicted frequency of serotiny (Fig. S5). The effect was greatest when examining the relationship between seed predation rate and the frequency of serotiny; a doubling of the mortality rate from 0.005 to 0.01 resulted in very low (<20%) serotiny at all squirrel densities. Increased mortality in adult stands results in increased turnover, allowing nonserotinous trees to replace dead trees more rapidly and resulting in unfavorable conditions for serotiny. We used mortality rates from ref. 11, which reported an increase in mortality rates for the genus *Pinus* in western North America since 1960 from ~0.0025 to 0.02. It is likely that contemporary patterns in serotiny reflect more the historical mortality rate than rates that have increased over the past several decades. Thus, our default value may somewhat underestimate the relative fitness of serotinous trees in historical stands. However, if increased mortality rates in mature stands are the new norm in lodgepole pine forests, serotiny may become increasingly disfavored, and mortality rates (or, more generally, nonfire distur-

bance rates) may be an underappreciated factor influencing the evolutionary dynamics of serotiny.

Postfire Stand Density. The curve relating prefire serotiny to postfire stand density has two parameters: the immigration rate γ_0 and the slope γ_1 . The model was moderately sensitive to immigration rate, with an increase of 1 SE resulting in a decline in serotiny of ~15% (Fig. S6). Because the model assumes that immigration is only from nonserotinous trees, it is expected that increasing this rate will favor nonserotinous trees. In actual stands, pollen dispersal in nonfire years, quantitative inheritance and incomplete heritability (3), and postfire dispersal of serotinous seeds from surrounding stands are likely to contribute serotinous individuals as well. Incorporating these dynamics into the model would likely decrease the sensitivity of the model to immigration rate. The model was less sensitive to changes in the slope of the postfire stand density curve (Fig. S7), with changes in the frequency of serotiny of less than 5% per SE.

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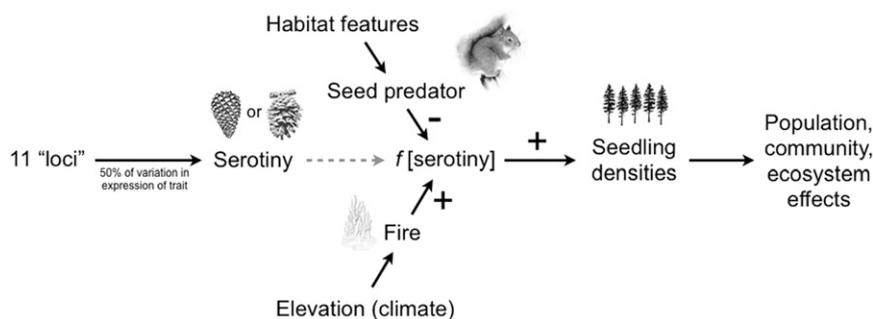


Fig. S1. Conceptual diagram of the interrelationships between genetic variation, selective agents, and ecosystem structure and function in lodgepole pine.

Table S1. Fixed effects from the generalized linear mixed model for cone survival

| Parameter | Estimate* | SE* | z | P |
|-------------------------------|--------------------|-------|-------|--------|
| Intercept | 1.90 | 0.59 | 3.20 | 0.001 |
| Cone type: serotinous (green) | -0.42 [†] | 0.074 | -5.71 | <0.001 |
| Cone type: serotinous (brown) | 1.48 [†] | 0.11 | 13.79 | <0.001 |
| Distance from midden | 0.27 | 0.29 | 0.92 | 0.36 |
| Distance × serotinous (green) | 0.19 | 0.086 | 2.19 | 0.028 |
| Distance × serotinous (brown) | 0.51 | 0.14 | 3.78 | <0.001 |
| DBH [‡] | 0.21 | 0.032 | 6.67 | <0.001 |

*Estimates and SEs are presented on the logit scale, standardized to mean = 0 and unit variance for numeric predictors.

[†]Parameters for cone type are relative to nonserotinous cones (i.e., the intercept).

[‡]Diameter at breast height, in centimeters.

Table S2. List of parameters and state variables

| Description | Symbol | Default | Range | SE | Source, ref. |
|-------------------------------|------------|---------|------------|-------|--------------|
| State variables | | | | | |
| Serotinous population size | n_s | | | | |
| Nonserotinous population size | n_n | | | | |
| Stand age | a | | | | |
| Selection parameters | | | | | |
| Fire return interval | F | n/a | 100–400 | | 1 |
| Squirrel density | S | n/a | 0–2.10 | | 2 |
| Other parameters | | | | | |
| Population growth rate | ρ | 0.05 | 0.01–0.21 | * | 3, 4 |
| Self-thinning: intercept | β_0 | 14.64 | | 0.24 | 5 |
| Self-thinning: slope | β_1 | -1.39 | | 0.055 | 5 |
| Adult mortality rate | μ | 0.005 | 0.005–0.02 | * | 6 |
| Immigration rate | γ_0 | 6.56 | | 0.21 | 7, 8 |
| Seedling density slope | γ_1 | 8.87 | | 0.37 | 7, 8 |

*No SE was available for these parameters, so the ranges were used for sensitivity analysis.

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Other Supporting Information Files

[Dataset S1 \(CSV\)](#)