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Climatic warming destabilizes forest ant communities

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How will ecological communities change in response to climate warming? Direct effects of temperature and indirect cascading effects of species interactions are already altering the structure of local communities, but the dynamics of community change are still poorly understood. We explore the cumulative effects of warming on the dynamics and turnover of forest ant communities that were warmed as part of a 5-year climate manipulation experiment at two sites in eastern North America. At the community level, warming consistently increased occupancy of nests and decreased extinction and nest abandonment. This consistency was largely driven by strong responses of a subset of thermophilic species at each site. As colonies of thermophilic species persisted in nests for longer periods of time under warmer temperatures, turnover was diminished, and species interactions were likely altered. We found that dynamical (Lyapunov) community stability decreased with warming both within and between sites. These results refute null expectations of simple temperature-driven increases in the activity and movement of thermophilic ecototherms. The reduction in stability under warming contrasts with the findings of previous studies that suggest resilience of species interactions to experimental and natural warming. In the face of warmer, no-analog climates, communities of the future may become increasingly fragile and unstable.

INTRODUCTION

Climate-driven shifts in community structure and function are already apparent in natural systems (1). Identifying the mechanisms that underlie these shifts in communities in response to climate change is vital to ecological forecasting efforts (2, 3). No species is an island, but many models of species responses to climate change make this assumption, ignoring indirect effects of climate change on species interactions. State-of-the-art forecasting models describe future communities at an equilibrium state determined by the joint effects of climate and species interactions (4, 5). However, this equilibrium perspective overlooks the dynamics of colonization, extinction, and patch occupancy of communities as they experience climate change. The outcomes of species interactions and community dynamics can allow us to forecast community resilience or decline under environmental perturbation, but only a limited number of statistical and theoretical models have addressed this question of system dynamics and stability under climate change (6–8). We still lack an understanding of how these processes operate in nature.

We explored the temporal dynamics and mathematical stability of the responses of ant communities to climate warming in one of the largest, geographically replicated experimental warming arrays in the world (9). The arrays were established in a warm, southern site (Duke Forest, located in the Piedmont region of North Carolina, USA) and a cool, northern site (Harvard Forest, located in the New England upland region of Massachusetts, USA). These sites are located, respectively, near the center and the northern range boundary of the deciduous forest biome in eastern North America. The experimental arrays spanned 6.5 degrees of latitude, yielding a mean annual temperature (MAT) difference of 5.8°C (Fig. 1A). Each site contained 15 open-top chambers, each 5 m in diameter, and warmed the forest floor year-round with thermostat-controlled forced air passed over hydronic heaters.

At each site, nine chambers were warmed in increments of 0.5°C from 1.5° to 5.5°C above ambient temperature; three additional chambers had forced air, but no heat, and a final three chamberless controls had no forced air and no heat (Fig. 1, A and B). The open-top chambers were also open at the bottom to allow free access to ants and other invertebrates. The range of temperatures spanned by the warming treatments encompasses climate projections of increased MAT from 1° to greater than 5°C over the next century (10). The warming increments represent a continuous experimental gradient of increasing temperature between sites: The unheated control chambers at the southern Duke Forest site had similar temperatures to the warmest heated chambers at the northern Harvard Forest site.

During the 5-year study, more than 60 species of ground-foraging ants were collected in the chambers, 30 of which occurred at both sites. Because most of these species forage a distance of less than 1 m from their nesting sites (11), the responses of ants to climate warming in these 5-m-diameter chambers primarily reflect the activities of resident colonies within the chambers, rather than transient movements of foragers or queens nesting outside of the chambers. More than 98% of foraging activity observed in the chambers involved workers originating from nests within the chambers (11). For ant foragers, with a typical body length of ~0.5 cm, the equivalent area of each chamber scales up to ~314 ha (1.2 square miles) in terms of human body lengths.

At the start of the experiment in 2010, we placed four artificial nest boxes into each of the 30 chambers; midway through the experiment, we added another four nest boxes. Because many ant species in eastern deciduous forests are nest-site limited, and competition for nest sites can be intense (12), artificial nest boxes are ideal experimental units for exploring the impacts of climate on interactions and associations among species. The boxes were constructed from carved wood blocks and were fitted with clear Plexiglas viewing windows that allowed us to census colonies with minimal disturbance (Fig. 1C). Censuses occurred approximately monthly during the growing season at each site over a span of 5 years of experimental warming (fig. S1 and table S1).

For the four or five most common species at each site (see Materials and Methods for species selection criteria), we pooled nest box census
records within each chamber and estimated the species-specific monthly binomial probability of colonization, extinction, and occupancy. Colonization and extinction were defined operationally as appearances and disappearances between consecutive censuses of a species in a nest box. Nest box occupancy per chamber ranged from 0.10 to 0.33 at Duke Forest and from 0.07 to 0.22 at Harvard Forest, with considerable dynamic turnover between consecutive monthly censuses and frequent replacement of one species by another with no intervening vacancy (table S1). Multiple occupancy of nest boxes was never observed: At each census, nest boxes were either occupied by a single colony or empty. These data suggest that nest boxes provided adequate nesting sites but were still sufficiently limited to reflect the dynamics of interactions and associations among species. Note that periodic nest box censuses provide indirect evidence of interactions among species through shifts in nest box usage as opposed to direct observations of interactions. Here and elsewhere, we refer to altered species interactions under climate change in a general sense, but because we acknowledge the possibility (albeit one that we consider unlikely) that some or all of the shifts in nest box usage in our experiment may be simple associations among species absent any interactions, we refer to the results from our experiment as shifts in species associations (see Results and Discussion for a full discussion of the nature of species interactions and associations in our system).

RESULTS AND DISCUSSION

We used multiple regression models with a quasi-binomial error structure [generalized linear models (GLMs)] to tease apart the effects of MAT in each chamber and the presence of other species on responses to warming. There were few cases in which colonization, occupancy, or extinction was affected by only MAT, as is often assumed for many species distribution models. Instead, the responses of species depended on both temperature and other species. Most species were connected to one or two other species in simple networks of positive and negative associations mediated by temperature (Fig. 2 and table S2). The responses of individual species to warming and to the presence of other species were complex and idiosyncratic, and these responses were different for colonization, extinction, and occupancy (Fig. 2).

Individual species responses to warming at Duke Forest
At Duke Forest, C. lineolata and, to a lesser degree, Aphaenogaster spp. were the most strongly influenced by the direct effects of warming. With
increasing warming, the probability of nest box occupancy by *C. lineolata* significantly increased and the probability of extinction decreased (figs. S2 to S4 and table S3). *Aphaenogaster* spp. exhibited decreased occupancy and colonization with increasing temperature (figs. S2 and S3 and table S3). For the remaining two focal species, *Brachyponera chinensis* and *Temnothorax curvispinosus*, occupancy, colonization, and extinction did not respond significantly to increasing chamber temperature. We found some cases in which occupancy of a focal species responded to temperature or other species, but we did not detect effects on colonization or extinction, possibly because of limited sample sizes (figs. S3 and S4 and table S3).

Although there is good evidence for the direct effects of temperature on *C. lineolata* and *Aphaenogaster* spp., there were also indirect effects of temperature on nest box occupancy, colonization, and extinction that were mediated by its effects on co-occurring species (fig. 2 and table S2). For example, the presence of *C. lineolata* was negatively associated with *Aphaenogaster* spp. occupancy and colonization; the presence of *B. chinensis* (a newly arrived exotic species at Duke Forest) was negatively associated with *C. lineolata* occupancy, and there was a significant interaction of MAT and *T. curvispinosus* occupancy on *C. lineolata* colonization.

**Individual species responses to warming at Harvard Forest**

At Harvard Forest, *Camponotus* spp., *Myrmica* spp., and *Temnothorax longispinosus* were the most strongly influenced by the direct effects of warming. Nest box occupancy and colonization of *Myrmica* spp. decreased with warming (figs. S5 and S6 and table S4), whereas occupancy by *Camponotus* spp. and *T. longispinosus* (fig. S5 and table S4) and colonization by *T. longispinosus* (fig. S6 and table S4) increased with warming. *Aphaenogaster* spp. and *Lasius* spp. did not exhibit significant occupancy or colonization responses to increasing chamber temperature. None of the five focal species exhibited significant effects of warming on the probability of extinction (fig. S7 and table S4).

We again found the widespread importance of the indirect effects of other species on occupancy, colonization, and extinction at Harvard Forest (fig. 2 and table S2). *Aphaenogaster* spp. and *Myrmica* spp. were each negatively associated with the other’s colonization, and *Myrmica* spp. and *Camponotus* spp. were each negatively associated with the other’s occupancy. The presence of *Camponotus* spp. and *Myrmica* spp. was negatively associated with *Aphaenogaster* spp. extinction and occupancy, respectively.

**Community responses to warming at Duke and Harvard forests**

In contrast to the idiosyncratic responses of individual species, responses at the community level were more consistent: Within and between sites, warming increased occupancy and decreased extinction, with non-significant, but positive, effects on colonization (fig. 3 and Table 1). There was no evidence that MAT influenced occupancy, colonization, or extinction differently between sites (no significant site × temperature interactions; table S5). When the sites were analyzed separately, occupancy increased and extinction decreased with temperature at Duke Forest. At Harvard Forest, the effects were marginally nonsignificant, but the slopes were of the same magnitude and direction as at Duke Forest (Table 1).

Community-level responses were driven by a few thermophilic species at each site that increased in occupancy and colonization in response to warming: *Camponotus* spp. and *T. longispinosus* at Harvard Forest and *C. lineolata* at Duke Forest. *Camponotus* spp., *C. lineolata*, and *T. longispinosus* have upper thermal tolerances between 42º and 46ºC, compared to upper thermal tolerances of 38º to 43ºC for the rest of the community (13). Because individual species responses were predictable on the basis of their upper thermal tolerances, community-level responses to warming were consistent.

**Community stability under warming**

To explicitly model the associations among species and link the outcomes of these associations to warming, we constructed a Markov transition matrix model (14, 15) for the data from each individual chamber. For a community of *n* interacting species, we constructed an *(n + 1) × (n + 1)* transition matrix, which includes a state for each species and one state for an empty patch. We recorded a total of 6960 individual state transitions in the 240 nest boxes and used these to construct 30 independent transition matrices, one for each chamber in the experimental array. The transition matrix describes the discrete-time dynamic occupancy of a patch based on the colonization, persistence, extinction, and turnover of individual species and their responses to each other. The model preserves the identity of each species and allows for asymmetries in the strength and outcome of species associations at each of the different chamber temperatures.

The left eigenvector of the transition matrix represents the vector of relative abundances of species at equilibrium, and the damping ratio of the first two eigenvalues represents the Lyapunov stability and measures return time to equilibrium following a small perturbation (16). Our parameterization of the Markov model does not assume that communities have reached an equilibrium state, but we do assume that transition probabilities for each chamber, averaged over the entire collecting period, were drawn from a stationary distribution.
These Markov models predicted a consistent shift in equilibrium species composition through time, with a set of thermophilic “winner” species becoming more common and an approximately equal number of “loser” species becoming less common. At Duke Forest, the equilibrium frequency of the thermophilic C. lineolata increased with warming, whereas the equilibrium frequencies of Aphaenogaster spp., T. curvispinosus, and empty patches either decreased or exhibited little change (fig. S8 and table S6). These shifts reflect the change in transition probabilities in the individual matrices: The transition (empty→C. lineolata) increased at higher temperatures, whereas the transitions (C. lineolata→Aphaenogaster spp.) and (C. lineolata→T. curvispinosus) decreased at higher temperatures. At Harvard Forest, there was a nonsignificant trend at higher temperatures toward a greater equilibrium frequency of the relatively thermophilic T. longispinosus (fig. S9 and table S6). Individual transition probabilities as functions of chamber temperature supported this result: The transition (T. longispinosus→empty) trended negatively with increased warming (table S7). Analyses of individual transition probabilities reinforced the results of the species association GLMs (fig. 2), which revealed evidence of indirect effects of warming at Harvard Forest. Although the transition (Myrmica spp.→Myrmica spp.) significantly increased with warming, there was also an increase in the transition (Myrmica spp.→T. longispinosus) with warming. The change in species associations with increasing temperature was further reflected in a decrease in the transitions (Aphaenogaster spp.→Myrmica spp.) and (Myrmica spp.→Aphaenogaster spp.) with warming.

The net effect of these altered species associations was the consistent reduction in the damping ratio of the transition matrix under warmer temperatures (Fig. 3A). This ratio measures the relative speed at which a perturbed system returns to its equilibrium (Lyapunov stability). Collectively, MAT and additive effects of site accounted for 35% of the variation among chambers in the damping ratio (Table 1). Thus, as temperatures increased, community stability decreased and did so at both sites. The stability and occupancy patterns predicted by the Markov models were not affected by using different criteria for including species in the model or by incorporating the possibility of measurement error in the estimation of transition probabilities (see “Alternative
foraging activity by ant workers closely tracks seasonal and diurnal fluc-
tuations in temperature (11, 20). However, the dynamics of nest site 
occupancy by colonies refute this null hypothesis of simple temperature-
driven activity levels. At higher temperatures, nest occupancy increases 
(Fig. 3B) because of a strong reduction in nest site abandonment (Fig. 
3D). These results were qualitatively robust to several variations on the 
formulation of the Markov model, such as treating net boxes as unavail-
able for colonization when occupied by a nonfocal species, changing the 
number of focal species used for the analysis, and replacing estimated 
transition probabilities of 0.0 with small nonzero probabilities reflecting 
measurement error (see “Alternative demographic and transition 
matrix model specifications” in the Supplementary Materials).

Limitations of our approach and priorities for future research
Our field-based manipulation of environmental temperature yielded 
evidence consistent with altered species associations among forest ant 
species and a reduction of dynamical stability at the community level. 
However, there are some limitations of our current study that future 
research efforts could address. Here, we highlight three such areas for 
further development. First, although we were able to quantify the 
consequences of warming with respect to ant community stability, 
we did not directly observe interactions among species leading to nest 
displacement. Given the strong nest site limitation in forest ants (21) 
and our direct observations of competitive interactions among ant 
species in and near the warming chambers (22), altered species inter-
actions seem the most likely explanation for our results. An alternative 
explanation for our results is possible, though one that we consider 
unlikely given the above. Our models may simply be detecting associations 
among species, absent any interactions, in their responses to warming. 
Detailed assessments of the nature and magnitude of altered species 
interactions under climate change are clearly needed—a call that has now 
been echoed by many climate researchers [reviewed by Angert et al. (23)]. 
Irrespective of the distinction between species interactions and associa-
tions, the outcomes of the 5-year climate manipulation experiment are

demographic and transition matrix model specifications” in the Sup-
plementary Materials).

To understand the mechanisms that disrupt community stability 
with warming, we analyzed the relationship between the Lyapunov 
stability of a matrix and its individual transition elements. Although 
the Lyapunov exponent reflects nonadditive contributions of all 
the transition elements in the matrix (16), the best correlate of the Lyapunov 
stability is the sum of the diagonal transition elements, which measures 
the probability that the system remains unchanged from one time step 
to the next (resistance). The higher the probability of species persistence 
from one time period to the next, the lower the Lyapunov stability (table 
S8). At the southern site and in warm chambers, persistence of colonies 
was high, community resilience was low, and the return to equilibrium 
was slow. In contrast, at the northern site and in cool chambers, persist-
ence of colonies was low, assemblage resilience was high, and the return 
to equilibrium was fast. When not perturbed, the system exists in a state 
of dynamic turnover; however, when perturbed by increasing environmental 
temperature, thermophilic colonies remain in nest sites for longer periods 
of time, reducing the overall stability of the community. Our metric 
of stability, the damping ratio, was uncorrelated with sample size, both 
including and excluding the empty class. Therefore, results for the 
damping ratio were driven by the structure of the matrices themselves 
rather than by variation in sample sizes used to construct the transition 
matrices in different chambers.

Null expectations for temperature effects on communities
Understanding the responses of ectotherms to climate change is especially 
challenging because of feedback between organismal physiology and 
behavior. Because the behavior and activity of ectotherms strongly 
depend on ambient temperature (17, 18), standard sampling procedures, 
such as pitfall traps may passively accumulate more individuals, even 
if the only effect of elevated temperatures is the increased random 
movement of foragers (19). For example, at the Duke Forest site, 
foraging activity by ant workers closely tracks seasonal and diurnal fluc-

Table 1. Occupancy, colonization, extinction, and stability responses to warming. For occupancy, colonization, and extinction, slope estimates ± 1 SE are 
from GLMs using a quasi-binomial error structure to examine the effect of MAT on community-wide occupancy, colonization, and extinction. F ratios and P 
values indicate the statistical significance of chamber temperature or site. The pseudo-$r^2$ is calculated as 1 − (residual deviance / null deviance). For stability, 
slope estimates ± 1 SE are from simple linear regressions to examine the effect of MAT on stability (damping ratio) for each site considered separately and from 
an ANCOVA with separate intercepts for site and a common slope for MAT (the site × MAT interaction was not significant and was dropped from the final 
model).

<table>
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<th>Data subset</th>
<th>Response</th>
<th>Predictor</th>
<th>Estimate</th>
<th>SE</th>
<th>F</th>
<th>P</th>
<th>$r^2$</th>
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<td>7.75</td>
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<td>0.0678</td>
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clear: Warming alters species-specific occupancy, colonization, and extinction rates; shifts the outcomes of pairwise associations among species; and consistently decreases the overall stability of temperate forest ant communities at two widely separated sites in eastern North America.

Second, our warming chambers, although being some of the largest of their kind, are limited to inferences at the chamber scale. However, given the small size of ants relative to the 5-m diameter of the warming chambers, entire ant communities are encapsulated within the footprint of a single chamber (see Materials and Methods). Because our study identifies a pattern of reduced stability under warming and suggests a mechanism via altered species interactions, then all else being equal, we would expect the pattern and process to scale up, that is, that the altered species interactions and dynamics of ant community responses to warming at the chamber scale would reflect responses at broader spatial scales of entire forests. Yet, it is still possible that our chamber-level results for dynamical stability may not scale up beyond the chamber or site level to the regional level. A regional scale or larger test of shifts in dynamical community stability under experimental climate change does not seem feasible with the infrastructure needed for a manipulative warming experiment. Testing whether our chamber results extend to larger spatial scales could be possible using space-for-time substitution (24). Indeed, this approach would allow for tests of dynamical stability from cooler and warmer background climates—an especially relevant comparison because temperate and tropical communities may respond differently to increasing temperature (25). However, as many researchers have noted, inferences from warmer geographic locations can, but do not necessarily, reflect responses to warming per se [reviewed by Bellard et al. (26)]. In this regard, it is interesting that both our higher and lower latitude warming arrays, situated at 42.5°N and 36°N latitude, respectively, yielded similar results, with cooler and warmer forest ant communities each showing a reduction in community stability under experimental warming. Many studies point to the greater susceptibility of lower-latitude (especially tropical) habitats to climate change, though some models and empirical work suggest greater susceptibility at mid- and higher latitudes (27, 28). Broader taxonomic and spatial coverage of community stability is needed to understand the biogeographic patterns of community stability and responses to global temperature rise.

Third and finally, although the reduction in community stability under warming translates to longer return times to equilibrium following an environmental perturbation, such as a marked increase in temperature, the long-term biological significance of a reduction in dynamical community stability is unclear. With slower recovery times following perturbation, warmed communities may be more fragile and susceptible to extirpation with increasing temperature (29). An important next step for this work then is to link the reduction in stability with the demographic and fitness consequences for individual species and biodiversity at the community level. Indeed, our analyses of the abundances of individual ant species and community composition in the warming arrays are already suggestive of potential higher-level consequences of the loss of dynamical stability. Within the warming chamber arrays, we detected substantial shifts in individual ant species abundances (13) and modest, but significant, shifts in ant community composition with increasing chamber temperature (20); these shifts tended to favor thermophilic subsets of communities, in some cases at the expense of less heat-tolerant species (see Supplementary Materials for a more detailed discussion of community dynamics and compositional changes).

Consequences of altered species interactions in a warmer world

What will communities of the future look like in a warmer world? Before species migrate from distant latitudes in an effort to track a changing climate (30), there will be shifts in abundance among resident species. It is this shift in abundance that is forecast with the Markov transition models of nest box dynamics. Although there is a strong focus in the literature on forecasting patterns of biodiversity change with warming (31), the effects of climate change on community dynamics are not well understood. Here, we provided experimental evidence for the cascading effects of warming on individual species and community-wide colonization and extinction processes, species associations, and, ultimately, community stability. Many researchers have pointed to the difficulty of incorporating species interactions into models of responses to climate change (32, 33). The idiosyncratic effects of climate and other species on colonization, extinction, and occupancy rates of individual ant species (Fig. 2) certainly do not lend themselves to simple generalizations. Nevertheless, at the community level, there were consistent increases in occupancy and decreases in extinction (Fig. 3). The dominance of thermophilic species increased in warmer experimental chambers, and this reduced the stability of local ant communities across climatically diverse sites.

Although the loss of community stability under warming may seem an intuitive result, decreasing stability is surprising for at least two reasons. First, decreasing stability of intact communities provides an important alternative to widely anticipated scenarios of “regime shifts” under climate change, in which communities switch abruptly to a contrasting alternative stable state (34). Second, some previous studies have found evidence for resilience, not destabilization, of trophic, predator-prey, and competition-based species interactions in response to climate change over geological (35, 36) and contemporary (6, 37, 38) time scales. However, these studies have relied on theoretical models, correlative data, and nondynamical community stability approaches. In contrast, our direct measurements of extinction and colonization dynamics in a long-term controlled field experiment suggest that climate warming not only may change the composition of ecological communities but also may lead to greater instability and slower recovery from environmental perturbation.

MATERIALS AND METHODS

Warming chambers

We constructed two experimental warming arrays at the center and the northern range boundary of the temperate deciduous forest biome in the eastern United States. The higher latitude array was located within Harvard Forest, Petersham, Massachusetts, and the lower latitude array was located within Duke Forest, Hillsborough, North Carolina. Details of the chamber construction and operation are provided elsewhere (9), but briefly, each site contained 15 chambers (5 m in diameter) that warmed the forest floor. Nine of these chambers were warmed via forced air in a regression design from 1.5° to 5.5°C above ambient in increments of 0.5°C; three additional chambers had forced air, but under ambient conditions, and a final three chambers had no forced air (“chamberless control”) and remained under ambient conditions.

We computed the MAT for each of the chambers with forced air (12 per site). The MAT of the chamberless control plots was computed for the single set of sensors located outside the warming chambers, and this value was assigned as the MAT for all three chamberless control plots. Raw temperature data were recorded at hourly intervals via a
ground-based sensor network of thermistors in the chambers; MAT is the mean of all hourly temperatures for the years in which the nest box censuses occurred.

Censuses of nest boxes

We constructed nest boxes from a wood block of untreated pine (14 cm × 15 cm × 2 cm) and balsa wood; we routed a zigzag pattern into the top of the nest box and cut an entryway in the side of the box. The nest box was covered on top with Plexiglas and a ceramic tile. The tiles were lifted to census the ant colony visible through the box. The nest box was covered on top with Plexiglas and a ceramic tile. The tiles were lifted to census the ant colony visible through the box. The nest box was covered on top with Plexiglas and a ceramic tile. The tiles were lifted to census the ant colony visible through the box. The nest box was covered on top with Plexiglas and a ceramic tile. The tiles were lifted to census the ant colony visible through the box. The nest box was covered on top with Plexiglas and a ceramic tile. The tiles were lifted to census the ant colony visible through the box.

An initial group of four nest boxes (A to D) per chamber was deployed at the beginning of the warming experiment, representing census points 1 to 41 at Duke Forest and census points 1 to 19 at Harvard Forest. A second group of four nest boxes (E to H) per chamber was deployed midway through the warming experiment, representing census points 26 to 41 at Duke Forest and census points 8 to 19 at Harvard Forest.

The nest boxes were censused approximately monthly during the growing season at Duke and Harvard forests (fig. S1). Note that fewer censuses occurred at Harvard Forest because the snow-free growing season is much shorter there than at Duke Forest. The number of days between censuses at Duke Forest ranged from 12 to 154, with a mean of 38.7 and an SD of 23.9. The first census occurred on 7 February 2011, and the last census occurred on 19 May 2015. The number of days between censuses at Harvard Forest ranged from 16 to 314, with a mean of 63.2 and an SD of 71.2. The first census occurred on 31 May 2012, and the last census occurred on 14 July 2015.

Species

Collectively, the two sites harbor 60 species of ground-foraging ants, with more than 30 species common to both sites. Ants make up more than half of macroinvertebrate abundance in eastern forests in North America, where they perform vital functions in ecosystems (39). Although some ant species in eastern U.S. forests frequently relocate nests (12), most species in this system foraged and dispersed over distances of less than 1 m (13). The foragers of resident colonies almost never ventured outside the chamber, and resident colonies usually remained within experimental chambers between consecutive censuses.

At Duke Forest, we found a total of 10 species (or species groups) inside the nest boxes: Stigmatomma pallipes (1 occurrence; 1 chamber), Aphaenogaster lamellidens (10 occurrences; 5 chambers), Aphaenogaster spp. (Aphaenogaster rudis and Aphaenogaster carolinensis) (444 occurrences; 15 chambers), B. chinesis (8 occurrences; 2 chambers), Camponotus castaneus (1 occurrence; 1 chamber), C. lineolata (466 occurrences; 15 chambers), Nyländeria flavipes (2 occurrences; 2 chambers), Prenolepis impar (1 occurrence; 1 chamber), Strumigenys spp. (1 occurrence; 1 chamber), and T. curvispinosus (79 occurrences; 9 chambers). From this group, we identified four focal species (or species groups) with sufficient replication for analysis and for which we are reasonably confident of their moving the entire colony into the nest box rather than their temporarily occupying the nest box as transient foragers: Aphaenogaster spp. (A. lamellidens combined with Aphaenogaster spp.; 454 occurrences; 15 chambers), B. chinesis (8 occurrences; 2 chambers), C. lineolata (466 occurrences; 15 chambers), and T. curvispinosus (79 occurrences; 9 chambers). At Harvard Forest, we found a total of 14 species (or species groups) inside the nest boxes: Aphaenogaster picea (62 occurrences; 15 chambers), Aphaenogaster fulva (9 occurrences; 7 chambers), Aphaenogaster spp. (92 occurrences; 15 chambers), Camponotus nearticus (1 occurrence; 1 chamber), Camponotus novaeboracensis (13 occurrences; 3 chambers), Camponotus pennsylvanicus (46 occurrences; 5 chambers), Formica spp. (1 occurrence; 1 chamber), Lasius alienus (8 occurrences; 4 chambers), Lasius nearticus (1 occurrence; 1 chamber), Lasius spp. (1 occurrence; 1 chamber), Myrmica punctiventris (21 occurrences; 11 chambers), Myrmica spp. (151 occurrences; 12 chambers), Tapinoma sessile (2 occurrences; 2 chambers), and T. longispinosus (25 occurrences; 6 chambers). From this group, we identified five focal species (or species groups) with sufficient replication for analysis and for which we are reasonably confident of their moving the entire colony into the nest box rather than their temporarily occupying the nest box as transient foragers: Aphaenogaster spp. (A. picea, A. rudis, and Aphaenogaster spp. combined; 163 occurrences; 15 chambers), Camponotus spp. (C. pennsylvanicus, C. nearticus, and C. novaeboracensis combined; 60 occurrences; 7 chambers), Lasius spp. (L. alienus, L. nearticus, and Lasius spp. combined; 10 occurrences; 5 chambers), Myrmica spp. (M. punctiventris and Myrmica spp. combined; 172 occurrences; 12 chambers), and T. longispinosus (25 occurrences; 6 chambers).

The ant species that we used to model the direct and indirect effects of climatic warming are a relatively small subset of the ant species present at each site and represent abundant ant species that typically nest in logs or small cavities and were thus amenable to living in nest boxes. Ant species with other nesting strategies (for example, soil nesting or arboreal nesting) and rare ant species were either too difficult to monitor or too data-poor for us to model their responses to warming. We note that although our chambers effectively contained entire ant communities, because nest box sampling plus alternative sampling methods, that is, pitfall trapping and litter sifting (9), yielded species diversity estimates on the order of regional levels of ant biodiversity (20), we focused our analyses on a subset of nest box–inhabiting species for practical reasons surrounding assessment and modeling.

Chamber 5 at Harvard Forest (a target warming increase of +5°C above ambient, with a MAT of 13.8°C) was too sparsely populated to estimate transition probabilities for our Markov model; thus, we eliminated chamber 5 from the Markov model and all other analyses. Chambers 3 and 10 at Harvard Forest (target warming increases of +2°C and +4.5°C above ambient with MATs of 10.7°C and 13.0°C, respectively) had estimated equilibrium frequencies of 1 for Camponotus spp.; thus, we added counts of 1 to the Camponotus transition matrix to allow for a small transition probability to the other states. Our results were qualitatively similar when we removed Camponotus entirely from these matrices or excluded chambers 3 and 10 from our analyses. For both sites, we used the criterion that a species (or species) group must be present during greater than one census point in at least two chambers to be included in the analyses.

Demographic response variables

We calculated proportions for the following: occupancy, as the number of times the focal species was present out of the total number of times the nest box was censused; colonization, as the number of times the nest box shifted from another state to the focal species out of the total number of times the species was present; extinction, as the number of times the species was present; and persistence, as the number of times the nest box contained the focal species and persisted with the focal species to the next census point.

We compared our results for occupancy calculated as the number of times the focal species was present out of the total number of times the nest box was censused to occupancy estimated from colonization
and extinction probabilities as the equilibrium frequency in the island-mainland variation of Levins’ (40) metapopulation model

\[ (\text{Occupancy}) = \frac{p(\text{Colonization})}{p(\text{Colonization}) + p(\text{Extinction})} \]

We further used colonization and extinction probabilities to estimate turnover

\[ (\text{Turnover}) = \frac{p(\text{Colonization}) \times p(\text{Extinction})}{p(\text{Colonization}) + p(\text{Extinction})} \]

**Statistical analysis**

All statistical models were performed using R version 3.2.2, *Fire Safety* (41). For each of the focal species at each site, we fit models of the proportion of nest boxes that were occupied, those that were colonized, and those that went extinct as functions of chamber MAT. Owing to issues with dispersion not equal to 1, we used GLMs with a quasi-binomial error structure. We conducted similar quasi-binomial GLMs for community-wide occupancy, colonization, and extinction at Duke Forest and Harvard Forest separately. We also fit models of the proportion of nest boxes that were inhabited by the focal species in one census point and persisted to the next census point. Like the occupancy, colonization, and extinction models, we used quasi-binomial GLMs to examine the impact of MAT of the chambers on persistence. We used *F* tests to assess the statistical significance of chamber temperature, because these tests are most appropriate for models where dispersion is estimated by moments (42).

We explored the effects of nonfocal species presence or absence over the course of the census period on each of the focal species responses (occupancy, colonization, and extinction) at each site. For each focal species and response variable, we constructed four models: (i) a main effect of chamber MAT, (ii) a main effect of nonfocal species presence or absence, (iii) both main effects of MAT and nonfocal species presence or absence, and (iv) both main effects plus the interaction of MAT and nonfocal species. We chose the best-fitting model among these four on the basis of the lowest Akaike information criteria (AIC) score (43). To obtain AIC values, we initially fit all models with a binomial error structure; we then refit the best-fitting models using a quasi-binomial error structure to obtain the correct SEs for the estimates.

We constructed a transition matrix for each chamber at each site, pooling observations across nest boxes within a chamber and transforming the raw counts to probabilities, such that columns summed to 1 (see table S1 for the raw number of transitions per chamber). We then used the *eigenanalysis* function from the *popbio* library (44) to perform eigenvalue decomposition on the probability matrix, allowing us to compute the Lyapunov stability (that is, the damping ratio, which is the ratio of the dominant eigenvalue to the second largest eigenvalue), the equilibrium frequencies for each species and the empty nest box class, and the transition probabilities for individual species and the empty nest box class.

Because the transition probabilities for each chamber are simple constants that do not incorporate density dependence, the Markov transition model does not forecast species extinctions. However, it does predict an equilibrium vector of relative abundances of each species (and of empty patches) that reflects all possible pairwise transitions in the system. None of the diagonal elements in the empirical matrices equaled 1.0; thus, there were no absorbing states. Instead, there is only a single equilibrium vector of relative abundances for each chamber, and the Lyapunov stability measures the return time to that equilibrium state following a small perturbation to the equilibrium. This Markov model effectively captures the interplay between occupancy, colonization, and extinction because the probabilistic transitions are always conditioned on the current state of the system.

To quantify which attributes of the transition matrices were most strongly contributing to the damping ratios, we examined Spearman’s rank correlation between damping ratio and the following: disturbance (the sum of the first row of the transition matrix that measures the tendency of the system to the empty state), evenness (the variance of the sums of the columns of the transition matrix), and resistance (the sum of the diagonals of the transition matrix that measures the tendency of the system to stay unchanged). We also calculated the return time, the number of time steps until the constant occupancy fraction is achieved under a simulation of 10,000 nest boxes starting in the empty state; we explored whether this alternative metric of stability was correlated with the damping ratio.

**SUPPLEMENTARY MATERIALS**

Supplementary material for this article is available at http://advances.sciencemag.org/cgi/content/full/2/10/e1600842/DC1

Levins’ metapopulation, turnover, and persistence models

Alternative demographic and transition matrix model specifications

Linking altered community dynamics with changes in community composition

**REFERENCES AND NOTES**


Climatic warming destabilizes forest ant communities
Sarah E. Diamond, Lauren M. Nichols, Shannon L. Pelini, Clint A. Penick, Grace W. Barber, Sara Helms Cahan, Robert R. Dunn, Aaron M. Ellison, Nathan J. Sanders and Nicholas J. Gotelli
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