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Weeds, worms, and deer: positive relationships among common forest understory stressors

Nicholas A. Fisichelli \bigcirc Kathryn M. Miller

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Abstract Biotic global change agents, such as nonnative plants ('weeds'), non-native earthworms ('worms'), and overabundant herbivores (white-tailed 'deer'), can be major stressors in the forest understory. The status and relationships among these global change stressors across large spatial extents and under naturally varying conditions are poorly understood. Here, through an observational study using a network of U.S. National Park Service forest health monitoring plots ($n = 350$) from eight parks in seven northeastern states, we modeled causal pathways among global change stressors through model selection in a structural equation (SEM) framework. Weeds, worms, and, deer were common across all parks in the study—46% of plots had non-native plants, 42% of plots had evidence of earthworms, and all parks had plots with high deer browse damage. All biotic global change stressors were significantly and positively correlated with one another (all Spearman rank correlations

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N. A. Fisichelli (\boxtimes)

Forest Ecology Program, Schoodic Institute at Acadia National Park, Winter Harbor, ME 04693, USA e-mail: NFisichelli@SchoodicInstitute.org

K. M. Miller

Northeast Temperate Network, National Park Service, Bar Harbor, ME 04609, USA

 > 0.44). Consequently, 28% of plots had a combination of earthworms absent, low deer browse, and no non-native plants, and 29% of plots included earthworms, non-native plants, and moderate or greater browse damage. Through SEM, we found strong support for pathways among global change stressors, e.g., deer browse positively influenced earthworm presence and both deer and earthworms promoted nonnative plants. Warmer air temperatures and higher soil pH also facilitated non-natives. This research highlights the tremendous multipronged management challenge for areas already experiencing the combined effects of weeds, worms, and deer and the future vulnerability of other areas as temperatures warm and conditions become more amenable to biotic global change stressors.

Keywords Climate - Earthworms - Forest soils - Herbivory - Global change - Non-native plants

Introduction

Global change stressors have far reaching effects across Earth's ecosystems. In addition to physical changes such as increasing temperatures, pollution, and habitat fragmentation, biotic changes in the form of introduced non-native plants ('weeds') and animals (such as earthworms, 'worms') and overabundant native herbivores (white-tailed 'deer') are major ecosystem drivers (Mack et al. [2000;](#page-10-0) Craven et al. 2016 ; Côté et al. 2004). These stressors are present across ecosystem types and protected area designations; management and restoration actions must take into account these multiple, interacting stressors for effective and realistic stewardship (Hobbs et al. [2014](#page-10-0)). For example, the northeastern U.S. includes national parks established for both significant natural features and to preserve cultural resources such as historic battlefields and military forts. In most cases, these historic parks were not originally designated to protect native biodiversity; yet, these cultural parks, with large areas of native ecosystems, have become important components of the protected area network in the heavily-populated and habitat-fragmented eastern U.S. As these parks look to management under continuously changing climate and related conditions in the near and long term (National Park System Advisory Board (NPSAB) [2012\)](#page-11-0), a major management strategy is to reduce existing stressors such as non-native species and overabundant herbivores (National Fish, Wildlife and Plants Climate Adaptation Partnership (NFWPCAP) [2012](#page-10-0)). Here, we take advantage of a network of forest monitoring plots in seven northeastern states to assess the status and relationships among non-native plants, non-native earthworms, deer browse pressure, and environmental conditions.

Individually, biotic stressors can have strong impacts on native plant performance, biodiversity, and ecosystem structure and function (Rooney and Waller [2003](#page-11-0); Vila` et al. [2011](#page-11-0); Craven et al. [2016](#page-10-0)). For example, chronic herbivory by white-tailed deer (Odocoileus virginianus), a native yet often overabundant species, simplifies understory community structure and can lead to decreases in native biodiversity (Rooney and Waller [2003;](#page-11-0) Côté et al. [2004\)](#page-10-0). Nonnative earthworms are ecosystem engineers, altering the forest floor and soil environment (Frelich et al. [2006;](#page-10-0) Suárez et al. [2006\)](#page-11-0) and reducing native seedling growth and survival (Dobson and Blossey [2015](#page-10-0); Dávalos et al. [2015a](#page-10-0)). Non-native plant invasions have myriad effects to native flora, including reductions in species diversity, abundance, growth, and fecundity (Mack et al. [2000;](#page-10-0) Vilà et al. [2011](#page-11-0)).

Increasingly, studies are finding positive and reinforcing relationships among biotic global change stressors (Nuzzo et al. [2009\)](#page-11-0). Overabundant deer facilitate increases in non-native plant species cover (Waller and Maas [2013](#page-11-0)) and may be required for successful non-native plant invasion and persistence (Kalisz et al. [2014\)](#page-10-0). Similarly, non-native earthworms promote non-native plant invasion (Bartuszevige et al. [2007;](#page-9-0) Roth et al. [2015;](#page-11-0) Craven et al. [2016](#page-10-0)). In combination, deer and earthworms change the forest floor environment, including leaf litter depth, plant cover and competition, and nutrient availability. These changes together negatively affect native seedlings (Dobson and Blossey [2015\)](#page-10-0) and promote establishment and growth of non-native plants (Dávalos et al. [2015b;](#page-10-0) Dobson and Blossey [2015\)](#page-10-0). Furthermore, overabundant deer appear to have a facilitative effect on invasive earthworms based on reductions in earthworms after deer exclosures are erected (Rearick et al. [2011](#page-11-0); Dávalos et al. [2015c](#page-10-0)), likely through changes to soil nutrient inputs, litter quality, and microbial communities (Karberg and Lilleskov [2009](#page-10-0)). Non-native plants may also have some reciprocal effects to promote deer and earthworms (Kourtev et al. [1999;](#page-10-0) Madritch and Lindroth [2009\)](#page-10-0).

Our expanding understanding of the interrelationships among non-native plants, earthworms, and deer come primarily from designed experiments in field and greenhouse settings, where treatments are applied and controlled (e.g., deer exclosures, removal of invasive plants, and addition of earthworms to microcosms). In this research, we ask whether relationships among biotic global change stressors are detectable in an observational study across a large region using naturally varying conditions, and further, whether our understanding of these relationships among biotic stressors allows us to model the system through structural equations. We hypothesize that deer have a positive effect on earthworm presence and that both deer and earthworms promote non-native plant species richness and cover. Understanding the status and relationships among above- and below-ground global change stressors is critical to land stewardship prioritization and efficacy.

Methods

Study sites

This study included eight national parks in seven states across the Northeast Temperate Network of the National Park Service (NPS) Inventory and Monitoring Program (Fig. 1). Parks range in size from 28 ha (Weir Farm National Historic Site, CT) to 14,577 ha (Acadia National Park, ME). These parks protect a diversity of natural and cultural heritage. For example, Roosevelt-Vanderbilt National Historic Sites (NY) preserve a country palace and the estate and homes of a former First Lady and President. Marsh-Billings-Rockefeller National Historical Park (VT) preserves the evolving history of conservation and forest management in the U.S. Acadia National Park protects the iconic coast, forest, and mountains of coastal Maine. All of the parks experienced intensive land-use prior to preservation. Some parks primarily only experienced timber harvesting (e.g., Marsh-Billings-Rockefeller and Acadia) while others also experienced agriculture and development (e.g., Minute Man National Historical Park, MA).

All of these parks now have large areas of forest protected managed by the NPS. These forests are

Fig. 1 Locations of national park units included in the study: 1. Acadia National Park, 2. Marsh-Billings-Rockefeller National Historical Park, 3. Saint-Gaudens National Historic Site, 4. Saratoga National Historical Park, 5. Minute Man National Historical Park, 6. Roosevelt-Vanderbilt National Historic Sites, 7. Weir Farm National Historic Site, and 8. Morristown National Historical Park. Study parks are found in three ecological provinces: Laurentian Mixed Forest (light gray), Adirondack-New England Mixed Forest (medium gray), and Eastern Broadleaf Forest (Oceanic) Province (dark gray) (Cleland et al. [2007\)](#page-10-0)

similar to neighboring forests outside park boundaries with the exception that park forests are often older with more complex structure (Miller et al. [2016](#page-10-0)). The parks are located across three ecological provinces: Laurentian Mixed Forest, Adirondack-New England

Mixed Forest, and Eastern Broadleaf Forest (Oceanic) Provinces (Fig. 1). Forest types range from spruce– fir–northern hardwoods in the north (ME, NH, VT) to transition hardwoods-white pine-hemlock (MA, NY) and central hardwoods in the south (CT, NJ).

Study sites have warm, moist, temperate climates. Annual mean temperature ranges across sites from 6.6 to $11.0 \degree C$ and summer (June, July, August) mean monthly temperature from 18.7 to 22.2 \degree C. Annual precipitation is abundant across this landscape, 1087–1447 mm, and similarly summer precipitation varies from 296 to 378 mm. January mean minimum temperatures vary from -5.78 °C in the south to $- 13.3$ °C at northern, interiors sites. Climate data are 2001–2010 averages from PRISM Climate Group [\(2012](#page-11-0)).

Forest sampling

Forest plot data used in these analyses are from the standard monitoring effort of the NPS Inventory and Monitoring Program. See Tierney et al. ([2016\)](#page-11-0) for details on sampling methods. Data are from 350 plots in eight parks (Electronic Supplementary Material Table S1). Each plot was sampled one time during the 4-year period, 2012–2015. Approximately 25% of the plots, distributed across parks, were sampled in each of the 4 years of the study. Forest plots are 225 m^2 $(15 \times 15 \text{ m})$ in Acadia, ME and 400 m² (20 \times 20 m) in the other seven parks. The number of plots per park varies from 10 for the smallest site (Weir Farm) to 176 for the largest site (Acadia). Sampling intensity varies from one plot per 2 hectares of forest (Saint Gaudens, NH and Weir Farm) to one plot per 73 hectares of forest (Acadia). Plot location is based on a generalized random stratified sampling design generated within each park.

Multiple forest health metrics were sampled in each plot, including species and diameter at breast height (DBH) for all trees ≥ 10 cm within the plot. Nonnative plant richness and cover data were collected within eight $1-m^2$ quadrats in each plot. All non-native plant species in each quadrat were identified to species and the percent cover (using a 10 cover-class system)

of each species was estimated visually. In addition, we collected multiple stand and site metrics, such as assessing deer browse damage and inspecting soils for evidence of earthworm presence.

Estimating deer population size is difficult. Accurate and standardized measures across large regions, meaningful at plot-level spatial scales, are not available. We did not attempt to calculate deer density, as the more proximate estimate of impact to the understory is deer browse damage to woody stems (Morellet et al. [2007](#page-10-0)). Thus, each plot was scored on a deer browse index with four impact levels (1-low, 2-moderate, 3-high, 4-very high). Low impact plots had no observed browse damage and palatable woody and herbaceous species present in the understory. Moderate-impact plots had evident but uncommon browse; high-impact plots included common browse evidence and/or rare browse-preferred species and browseresilient vegetation limited in height growth by deer browsing. Very high impact plots had omnipresent browse evidence, a distinct browse line around 1.5 m above the ground, browse-preferred species were absent, and non-preferred species showed signs of heavy repeated browsing.

Field sampling crews assessed direct and indirect evidence of non-native earthworms in three different locations within each plot. This included casts, middens, earthworm burrows, and earthworms detected during soil sampling. This rapid technique is similar to other protocols (Loss et al. [2013](#page-10-0); Fisichelli et al. [2013\)](#page-10-0) and captures presence of nonnative earthworms but does not capture species composition or abundance.

In addition to biotic stressors, we included several environmental variables assessed at the plot level (Table 1). To quantify soil pH, the upper (A) mineral soil horizon was collected from each plot. In plots where the organic soil horizon was directly above bedrock (no mineral soil horizon present), a sample of the organic horizon was collected instead. Soil pH was measured in distilled water with an electronic pH meter. Organic layer depth was also measured. Climate data for each plot were extracted from PRISM 800 m² resolution climate surfaces (2001–2010 averages, PRISM Climate Group [2012](#page-11-0)).

Analyses

All analyses were conducted in R v3.3.2 (R Core Team [2016\)](#page-11-0). All data were averaged to the plot level, e.g., number of non-native species and percent cover from the eight $1-m^2$ quadrats. We calculated Spearman rank correlations (ρ) and significance tests among biotic global change drivers (deer browse, non-native earthworm presence, non-native plant richness, and nonnative plant cover) and among environmental variables. Spearman rank correlations are appropriate for non-normally distributed and ordinal data (Sokal and Rohlf [1995](#page-11-0)). Due to lack of complete independence between non-native plant richness and cover (cover $= 0$ whenever richness $= 0$), this correlation was only calculated for plots with at least one nonnative species present (160 of 350 plots).

To more holistically assess relationships among biotic stressors in the understory, we utilized generalized linear mixed-effects models, model comparison with Akaike information criteria (AIC), and a structural equation modeling (SEM) framework. SEM is a useful framework for gaining insights into the direct and indirect relationships among variables in ecological systems (Grace [2006](#page-10-0)). SEM is a method for inferring and testing causal relationships among variables and is most often used in observational studies (Grace et al. [2012](#page-10-0)). Recent advances have made this tool more flexible and useful for ecological data (Grace et al. [2012;](#page-10-0) Lefcheck [2015](#page-10-0)). Historically,

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SEM has required assumptions of normally distributed data and independence among observations (Grace [2006\)](#page-10-0). Direct acyclic or piecewise SEM allows for inclusion of response variables with non-normal distributions, random effects, and different correlation structures using local estimation (Lefcheck [2015\)](#page-10-0). In our study, we were able to include response variables with Gaussian, Poisson, and binomial error distributions and the hierarchical structure of the research design (blocks nested in site). Piecewise SEM supports confirmatory path analysis but has some limitations such as inability to evaluate feedback loops or include latent variables. For example, the feedback loop of deer to earthworms and back to deer cannot be assessed here and is likely better examined with longitudinal data. SEM analyses utilized the piecewise SEM package (Lefcheck [2015](#page-10-0)).

We used pre-existing knowledge about the system to construct a structural equation model and hypothesized causal relationships (pathways) among variables within the model (see Fig. S1 for full hypothesized model, key pathways highlighted below). We are interpreting the relationships uncovered in this observational study as quantitative causal claims based on the assumptions developed from previous experiments and that these causal relationships are likely transferable to other datasets from similar forest types in similar environmental conditions (Pearl [2012\)](#page-11-0). Statistical tests of the pathways in the SEM do not prove causation but rather provide support for the causal claims. In SEM, single-headed arrows represent direct causal relationships, with the variable at the tail of the arrow representing the predictor of the variable at the head of the arrow. The overarching goal was to test the following hypothesized relationships among biotic global change stressors (see ''[Introduction](#page-0-0)'' for justifications for these relationships):

Earthworm presence \leftarrow Deer browse Non-native plant richness \leftarrow Earthworm presence Non-native plant richness \leftarrow Deer browse Non-native plant cover \leftarrow Earthworm presence Non-native plant cover \leftarrow Deer browse Non-native plant cover \leftarrow Non-native plant richness

Two environmental drivers were included in the SEM, soil pH and summer mean monthly air temperature (JJA). Both variables are known to affect earthworms and non-native plants. Higher soil pH is more conducive to non-native plant and earthworm invasion than acidic, low-nutrient soils (Alpert et al. [2000;](#page-9-0) Fisichelli et al. [2013\)](#page-10-0). Warmer temperatures facilitate non-native species colonization, reproduction, and population persistence (Walther et al. [2009](#page-11-0)). Furthermore, warmer temperatures are related to nonnative plant species occurrence in forest plots in the region (Fisichelli et al. [2014](#page-10-0); Oswalt et al. [2015\)](#page-11-0). For earthworms, extremes of hot and cold temperatures can be limiting. Within the north temperate forests in this study, warmer temperatures likely promote earthworm occurrence (Curry [2004\)](#page-10-0). There are many other potential environmental variables that could have been included in analyses. We selected soil pH and temperature due to their known effects to response variables, straightforward measurement techniques, and known correlations with other drivers (Table S2). For example, summer temperature is positively related to summer precipitation ($\rho = 0.85$, $p < 0.001$). Soil pH is also positively correlated to summer precipitation ($\rho = 0.64$, $p < 0.001$) and negatively correlated with overstory conifer basal area ($\rho = -0.51$, $p < 0.001$), so higher soil pH plots received more summer precipitation and tended to have fewer conifers in the overstory. Thus, these individual environmental drivers most accurately reflect a suite of related understory conditions. Additional explanatory variables such as deer density were not available across all study parks and thus omitted from analyses. Here, we focus on the global change stressors, including deer browse pressure, and their relationships.

The SEM contains three component models, with the response variables earthworms (presence and absence), non-native species richness, and non-native plant cover. We developed generalized linear mixedeffects models with the associated response variable distribution (binomial for earthworm presence and absence, Poisson for non-native species richness, and Gaussian for non-native plant cover). For non-native plant cover, we only included plots with non-native plants present $(n = 160)$ since non-native plant absence was modeled in the non-native plant richness analyses and omission of the plots with 0% cover permitted a Gaussian approach. Each mixed-effects model included a random effect of park subunit $(n = 15)$ nested in park $(n = 8)$ (i.e., block nested in site) and used maximum likelihood estimation with the lme4 package (Bates et al. [2015](#page-10-0)). The non-native species richness model included an additional individual-level random effect because of overdispersion (Harrison [2014](#page-10-0)). Based on the hypothesized SEM, we included three explanatory variables for earthworm presence (deer browse, summer temperature, and soil pH) in the full model. For non-native plant richness we included four explanatory variables (deer browse, earthworm presence, summer temperature, and soil pH), and for non-native plant cover we used five explanatory variables (deer browse, earthworm presence, non-native plant richness, summer temperature, and soil pH). Non-native plant cover was natural log transformed to meet statistical assumptions (i.e., linear relationships and homoscedasticity of errors). We used bivariate scatterplots and residual plots to visually assess relationships, model fit, and normality of errors.

We conducted model comparison for each SEM component model using AIC (Burnham and Anderson [2002\)](#page-10-0). We compared models varying in complexity from inclusion of only a single explanatory variable to a model with all additive effects. Earthworm, nonnative plant richness, and non-native plant cover models with the lowest AIC values ($\Delta AIC < 2$) and fewest parameters are considered the best fit (Burnham and Anderson [2002\)](#page-10-0). We also calculated and report individual pathway unstandardized coefficient estimates, standard errors, and significance tests using the Kenward–Rogers approximation (Lefcheck [2015](#page-10-0)). It is not possible to calculate standardized path coefficients for generalized linear models with Poisson or binomially distributed response variables since these response variables cannot be standardized. We calculated variance explained (R^2) for each generalized linear mixed-effects model using the methods of Nakagawa and Schielzeth [\(2013](#page-10-0)) and report the marginal R^2 for the fixed effects included in the SEM.

Results

Biotic global change stressors were abundant across plots (Fig. [2;](#page-6-0) Table S3). Evidence of non-native earthworm presence was found at 42% of plots (147 of 350 plots). Earthworm presence was lowest at Acadia, ME (9% of plots) and highest at Morristown, NJ (100%) and Weir Farm, CT (100%). Deer browse

b Fig. 2 Bivariate relationships among biotic global change stressors (plot-level averages, $n = 350$). Spearman rank correlations are shown in the upper left corner of each subpanel (all correlation p values \lt 0.001). Deer browse damage increases from 1 (low) to 4 (very high). Non-native plant values are plot averages per m^2 . Boxplot boxes are the interquartile range, thick horizontal lines are the medians, dashed vertical lines are the outer tails (1.5 \times interquartile range), and open circles are plot values beyond the outer tails

damage was observed in all study parks (overall index mean $= 1.83$ on a scale from 1 to 4) and all parks had at least three plots with high browse damage (in $dex = 3$). Low browse damage (index = 1) was most common (44%, 153 plots) and very high damage $(index = 4)$ least common $(5\%, 17$ plots). Overall, the lowest browse pressure was found in Acadia [mean $(SD) = 1.4$ (0.6)], Marsh-Billings-Rockefeller, VT [mean $(SD) = 1.6$ (0.8)], and Saint-Gaudens, NH [mean $(SD) = 1.6 (0.8)$]. The highest mean browse pressure was found at Morristown [mean $(SD) = 3.2$] (0.7)] and Weir Farm [mean $(SD) = 2.9$ (1.0)]. Nonnative plants were present at all parks and in 46% of plots (160 plots) overall. Plots averaged 0.72 nonnative species $(SD = 1.32)$ per 1-m² subplot. Nonnative plant cover averaged 6.3% (SD = 16.4), with a maximum plot average of 122%. The most heavily invaded parks include Saratoga National Historical Park, NY [means (SD): 2.9 (2.1) species, 21 (23)% cover], Morristown [means (SD): 2.3 (1.6) species, 34 (35)% cover], and Minute Man, MA [means (SD): 1.8 (0.9) species, 17 (13)% cover]. The least invaded sites include Acadia [means (SD): 0.05 (0.2) species, 0.2 (1.1)% cover] and Marsh-Billings-Rockefeller [means (SD): 0.2 (0.3) species, 0.4 (0.8)% cover].

All biotic global change stressors were significantly correlated with one another (all Spearman rank correlations ≥ 0.44 and p values $\lt 0.001$) (Fig. 2). All bivariate plots show positive relationships among biotic stressors, thus earthworm presence increases with deer browse, and non-native plant richness and cover both increase with earthworm presence and deer browse. Consequently, 28% of plots (101 plots) had earthworms absent, low deer browse (index $= 1$) and no non-native plants. Furthermore, 29% of plots (103 plots) included earthworms, non-native plants, and moderate or more severe browse damage.

The hypothesized SEM fit the data well and supported multiple pathways among global change

stressors (Table [2;](#page-7-0) Fig. [3](#page-8-0); Table S4). All supported pathways show positive relationships among variables. Evidence of non-native earthworms was best fit (lowest AIC) with the full hypothesized model including all three explanatory variables (deer browse, soil pH, and summer temperature). The second best fit model omitted deer browse and had a $\Delta AIC = 7.4$. Non-native plant richness was best fit with the model containing soil pH and earthworms ($\Delta AIC = 0.3$ and fewest parameters [2] among models with $\Delta AIC\lt2$). There was also some support for the hypothesized pathway between summer temperature and non-native richness (path coefficient p value \lt 0.05 and inclusion in the lowest AIC model). The more complex, hypothesized model including four explanatory variables (soil pH, earthworms, deer, and temperature) performed similarly ($\triangle AIC$ 1.7) to the simpler twovariable model with only soil pH and earthworms. Similar to richness, non-native plant cover was best explained by a two-parameter model including deer browse and non-native plant richness ($\Delta AIC = 0.5$) and fewest parameters [2] among models with $\Delta AIC < 2$). There was marginal support for the pathway between soil pH and non-native plant cover (p value $\lt 0.1$ and inclusion in the lowest AIC model). The hypothesized plant cover model with five explanatory variables was not supported $(AAIC = 3.3)$. Component models explained 33–67% of the variance (R^2) , based on fixed effects.

The SEM supported direct effects of deer browse, soil pH, and air temperature on non-native plant measures as well as indirect positive effects mediated through earthworm presence and non-native plant richness (Fig. [3](#page-8-0)). For example, soil pH effects on both non-native plant variables occurred through direct pathways (positive effect) and indirectly through soil pH influence on earthworm presence (positive effect). Deer browse and air temperature showed similar direct and indirect relationships to non-native plant variables. Earthworm presence also directly influenced non-native plant richness and indirectly influenced cover through its effect on richness.

Discussion

Deer browse damage, non-native earthworms, and non-native plants are present within forest understories across the northeastern U.S., including within sites

Pathway	Estimate	SE.	<i>p</i> value
Earthworm presence \leftarrow Soil pH	1.614	0.267	< 0.0001
Earthworm presence \leftarrow Summer temperature	0.917	0.144	< 0.0001
Earthworm presence \leftarrow Deer browse	0.701	0.233	0.003
Non-native plant richness \leftarrow Soil pH	0.435	0.084	< 0.0001
Non-native plant richness \leftarrow Summer temperature	0.363	0.184	0.048
Non-native plant richness \leftarrow Deer browse	0.041	0.077	0.60
Non-native plant richness \leftarrow Earthworm presence	0.485	0.191	0.01
Non-native plant cover \leftarrow Soil pH	0.464	0.257	0.07
Non-native plant cover \leftarrow Summer temperature	0.156	0.382	0.684
Non-native plant cover \leftarrow Deer browse	0.390	0.203	0.058
Non-native plant cover \leftarrow Earthworm presence	0.253	0.437	0.56
Non-native plant cover \leftarrow Non-native plant richness	0.821	0.130	< 0.0001
Soil pH \leftrightarrow Summer temperature	0.537		< 0.0001

Table 2 Unstandardized path coefficients, standard errors, and p values for all hypothesized fixed-effects pathways in the full SEM (Fig. [3](#page-8-0))

Single-headed arrows point from predictor to response variable and two-headed arrows indicate unresolved correlations in the model. Bold black font denotes p values < 0.05 , italics font denotes marginally significant pathways (p value 0.05–0.10), and gray font pathways were not supported. Significance tests were based on Kenward–Rogers approximation (see "[Methods](#page-1-0)"). Three component models within the SEM are generalized linear mixed-effects models with the responses earthworm presence (binomial), non-native plant richness (Poisson), and (ln)non-native plant cover (Gaussian)

receiving the highest levels of protection (i.e., national parks). All these biotic global change stressors are positively correlated with one another; moreover, there was support for four out of six hypothesized relationships among these biotic stressors in the SEM. Environmental conditions, namely higher soil pH, warmer temperatures, and related factors, appear to contribute to the presence and magnitude of these biotic stressors. The variation in and strong relationships among understory stressors across the landscape suggest opportunities to preserve native forest conditions in areas not yet severely impacted as well as the tremendous multipronged management challenge for areas already experiencing the combined effects of weeds, worms, and deer.

Biotic global change agents

As mentioned above, most hypothesized relationships among biotic global change stressors, based on a large body of scientific knowledge (e.g., Côté et al. [2004](#page-10-0); Nuzzo et al. [2009](#page-11-0); Rearick et al. [2011](#page-11-0); Waller and Maas [2013](#page-11-0); Craven et al. [2016\)](#page-10-0), were supported in analyses. Thus, across a large number of sites and including plots in all available forest types, deer affect earthworms and deer and earthworms affect nonnative plants. These are the types of positive relationships that may lead to invasional meltdowns (Simberloff and Von Holle [1999](#page-11-0)). Reciprocal relationships among these variables likely also exist in specific circumstances, though we were unable to test for them due to the constraints of the piecewise SEM structure (Lefcheck [2015](#page-10-0)). For example, palatable non-native plant species with high quality leaf litter may promote both deer and earthworms, especially within local areas with otherwise unpalatable native species producing low-quality litter and acidic soils (Kourtev et al. [1999](#page-10-0)).

The presence and strengths of biotic global change stressors across large landscapes are better understood for non-native plants than earthworms or deer browse pressure. Patterns of non-native plants across parks in the study generally match nearby parks in the National Park System (Fisichelli et al. [2014](#page-10-0)) and USFS Forest Inventory and Analysis (FIA) data that includes plots under all ownership types in the study region (Oswalt et al. [2015\)](#page-11-0). This research captures overall non-native plant richness and cover, which both relate strongly to other biotic stressors and environmental conditions. Although we did not look at individual non-native

Fig. 3 Structural equation model (SEM) illustrating supported relationships among biotic global change stressors and environmental variables. Numbers on the arrows are unstandardized path coefficients. Bold black arrows are supported pathways in the best fit, parsimonious model $(\Delta AIC < 2$ and fewest parameters), medium gray arrows are additional pathways in the lowest AIC model ($\Delta AIC = 0$), and fine gray arrows are unsupported pathways (more complicated models with $\Delta AIC > 0$, coefficients not shown). Bold black pathway coefficients have a p value $\lt 0.05$ and gray coefficients are marginally significant (p value 0.05–0.10). Three component models within the SEM are generalized linear mixed-effects models with the responses: earthworm presence (binomial), non-native plant richness (Poisson), and (ln)non-native plant cover (Gaussian). Percentages are the variance explained by fixed effects for each response variable (marginal R^2)

plant species, the most abundant species in these parks include Berberis thunbergii, Frangula alnus, and Lonicera spp., all species known to be associated with deer and earthworm impacts (Kourtev et al. [1999](#page-10-0); Frelich et al. [2006\)](#page-10-0).

This study also provides evidence for ubiquitous earthworm presence across multiple states in the northeast U.S., similar to findings for mid-Atlantic states (Dávalos et al. $2015c$) and the western Great Lakes region (Loss et al. [2013;](#page-10-0) Fisichelli et al. [2013](#page-10-0)). We did not collect data on earthworm density, biomass, or species composition, although all have been shown to influence understory plants and soil conditions (Suárez et al. [2006;](#page-11-0) Craven et al. [2016](#page-10-0)). Non-native earthworm community characteristics have been documented in research at smaller spatial scales in the region, though abundances and species distributions remain uncertain at larger spatial scales (Hendrix and Bohlen [2002\)](#page-10-0).

Similar to earthworms, we found pervasive evidence of browse damage within plots at all study sites. Deer densities across eastern forests are considered chronically high (McShea [2012\)](#page-10-0), though we did not attempt to compare browse pressure with an historical range of conditions to quantitatively assess whether the influence of deer can be considered extreme and a global change stressor. Browse damage is a more proximal measure of deer influence on vegetation than deer population estimates but may not fully capture variation in browse preference and vegetation availability within the browse layer (0.2–2 m above ground level). Regardless, selective herbivory is a major driver of understory plant performance (Fisichelli et al. [2012\)](#page-10-0) and the high browse levels found in many plots and the browse relationships to weeds and worms confirm the importance of this stressor.

This research included a single census (completed over 4 years) and thus is a snapshot in time and does not reflect biotic stressor temporal dynamics within the system. The rate of responses (e.g., increase in percent cover of non-native plants) and interannual variability are not known. However, deer browse damage to woody plants is detectable for many years and non-native plants in the study include perennial herbaceous and woody species likely detectable each year. Similarly, evidence of earthworms, including middens and removal of the organic layer are relatively persistent signs. Thus, although some variables may show some interannual variability (e.g., percent cover and number of non-native species), the relationships quantified here likely persist over time. The plots included in this study are permanent, longterm monitoring plots sampled on a 4-year cycle and thus temporal variability can be assessed in the future.

Environmental conditions

We selected a small number of environmental variables and recognize that these may reflect other important and unmeasured factors, such as are present in any observational study of a complex system. Soil pH was a predictor of all biotic stressors in the SEM. Soil pH directly influences the soil microbial community (Fierer and Jackson [2006](#page-10-0)) and the cycling and availability of nutrients and metals (e.g., aluminum) in the soil (Schoenholtz et al. [2000\)](#page-11-0), with cascading

effects on earthworms and non-native plants (Kourtev et al. [1999](#page-10-0); Frelich et al. [2006\)](#page-10-0). Similarly, more basic and nutrient rich soils also harbor larger numbers and cover of non-native species (Dávalos et al. [2015c](#page-10-0)). Soil pH reflects the current overstory (e.g., the conifer component and acidic needleleaf litter inputs), past vegetation composition, and historical land use (e.g., soil tilling) (Schoenholtz et al. [2000](#page-11-0)). Soil pH was also correlated with summer temperature; cooler northern sites were more acidic (and had a greater conifer overstory component). Thus, soil pH likely influenced the system through multiple effects and multiple related factors.

Similar to soil pH, temperature relates to multiple factors. Warmer sites in the study are to the south and are in generally more fragmented landscapes with greater residential and commercial development, agriculture, and transportation corridors (Theobald [2013\)](#page-11-0). These areas likely have had greater opportunity for human-mediated introduction of non-native plants and earthworms onto the landscape (Hendrix and Bohlen [2002](#page-10-0); Oswalt et al. [2015](#page-11-0)). Furthermore, fragmented landscapes with abundant edge habitat and adjacent agricultural fields can support larger numbers of deer than intact forests (Côté et al. [2004](#page-10-0)). Thus, landscape conditions and history likely promote biotic stressors at a broader spatial scale while plot level conditions (e.g., soil pH) determine local-level presence and severity of biotic stressors.

Restoration and adaptation

The chronic nature of browse damage in forest understories (McShea [2012](#page-10-0)) and linkages to earthworms, non-native plants, and other global change stressors such as temperature (Fisichelli et al. [2012\)](#page-10-0) indicate browse pressure as a key stressor and potential intervention point for successful management. Focusing effort solely on non-native plant eradication may not have desired outcomes if deer browse pressure and earthworms are not simultaneously addressed (Nuzzo et al. [2009;](#page-11-0) Waller and Maas [2013\)](#page-11-0). Although often challenging to implement due to social and political hurdles, reductions in deer populations are possible and can have desired effects (McShea [2012\)](#page-10-0). Unfortunately for non-native earthworm management, we are unaware of any feasible control measures to eradicate established populations.

Another global change agent, warming temperatures, is likely to affect forest understory dynamics and promote the biotic stressors (based on the SEM results). Cold, acidic, conifer dominated forests currently have relatively low stressor levels. All parks in the study have already experienced significant warming (Monahan and Fisichelli [2014\)](#page-10-0) and it is unclear how or if forest composition will change in the future as temperatures continue to warm. Northern conifer species present at parks in the study are projected to undergo substantial decreases in potential suitable habitat as temperatures warm, while at the same time broadleaf species gain suitable habitat (Fisichelli et al. [2014](#page-10-0)). Because deer, earthworms, and nonnative plants are already present at all parks, decline of the current overstory and shift in tree composition to broadleaf species could allow these biotic stressors to establish, assuming soil pH and other conditions are suitable. Early detection and rapid response to control these stressors may be effective in these areas where they are not yet widely established (Abella 2014). It is important to note that responses by native biota are species-specific and some may perform well in the presence of non-native plants (Dávalos et al. [2015a](#page-10-0)), benefit from browse pressure on competing palatable plants (Rooney and Waller [2003](#page-11-0)), or respond positively to leaf litter loss, such as from earthworm consumption (Warren et al. [2012\)](#page-11-0). Understanding the status and relationships among understory global change stressors and associated environmental variables is critical to identifying vulnerable sites and developing successful management strategies tailored to individual site conditions and priority native species.

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