Precipitating impacts of climate change on habitat connectivity of *Kalopanax septemlobus* in South Korea

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**A R T I C L E   I N F O**

**Article history:**
Received 24 July 2015
Received in revised form 2 January 2016
Accepted 11 January 2016
Available online 21 January 2016

**Keywords:**
Castor aralia 
Graph theory 
Habitat suitability 
Maximum entropy modeling 
Probability of connectivity

**A B S T R A C T**

Understanding the drivers of habitat distribution patterns and assessing habitat connectivity are crucial for conservation in the face of climate change. In this study, we examined a sparsely distributed tree species, *Kalopanax septemlobus* (Araliaceae), which has been heavily disturbed by human use in temperate forests of South Korea. We used maximum entropy distribution modeling (MaxEnt) to identify the climatic and topographic factors driving the distribution of the species. Then, we constructed habitat models under current and projected climate conditions for the year 2050 and evaluated changes in the extent and connectivity of the *K. septemlobus* habitat. Annual mean temperature and terrain slope were the two most important predictors of species distribution. Our models predicted the range shift of *K. septemlobus* toward higher elevations under medium-low and high emissions scenarios for 2050, with dramatic reductions in suitable habitat (51% and 85%, respectively). In addition, connectivity analysis indicated that climate change is expected to reduce future levels of habitat connectivity. Even under the Representative Construction Pathway (RCP) 4.5 medium-low warming scenario, the projected climate conditions will decrease habitat connectivity by 78%. Overall, suitable habitats for *K. septemlobus* populations will likely become more isolated depending on the severity of global warming. The approach presented here can be used to efficiently assess species and habitat vulnerability to climate change.

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1. **Introduction**

Climate change is regarded as one of the major drivers of changes in biodiversity and ecosystems, and its possible effects are receiving worldwide attention (Duraiappah et al., 2005; Sala et al., 2000). Indications for impacts of climate change have already been found in many species over a wide taxonomic range (Nussey et al., 2005; Parmesan, 2006; Parmesan and Yohe, 2003). In addition, such impacts may be reinforced by habitat loss and fragmentation, which can lead to species range shifts and a reduction in habitat connectivity (Honnay et al., 2002; Opdam and Wascher, 2004).

Although species may have the capacity to shift their range, long distances between habitats or other barriers may restrict their movement. Moreover, the ability to move across the landscape depends on species-specific behavior and landscape structure (Béislé, 2005; Goodwin and Fahrig, 2002). Thus, understanding the range-shifting capacities of species in response to climate change has important conservation implications for the predictions of future extinction risk and distribution changes (Angert et al., 2011). More importantly, because climate change appears to be inevitable, an effective adaptation strategy may involve preserving and restoring landscape connectivity for long-term persistence of ecological processes, such as dispersal and gene flow (Crooks and Sanjayan, 2006; Rosenberg et al., 1997; Templeton et al., 2001).

Landscape connectivity, or the degree to which the landscape facilitates or disturbs movement among resource patches (Taylor et al., 1993), affects dispersal success and colonization rates (With and King, 1999a, 1999b). Such patterns in turn influence the biodiversity, ecosystem function, and resilience of species to climate change (Fahrig, 2003; Gonzalez et al., 2009; Lawler, 2009). Specifically, well-connected landscapes may enable tracking of species’ suitable climate and habitat conditions through time and
thereby may allow ecological and evolutionary processes to be sustained. Increasingly, ecologists have reported that improving connectivity is necessary for biodiversity conservation and is thus one of the most commonly recommended strategies for helping species adapt quickly and survive rapid climate change (Heller and Zavaleta, 2009; Theobald et al., 2012).

Even though global warming has begun to affect the regional climate system (Boo et al., 2004; Jung et al., 2002) and cause range shifts of terrestrial flora in many countries (Allen et al., 2010), it is unknown whether existing biosphere reserves and semi-natural forests may ensure species’ long-term persistence. In this study, we use a species distribution modeling tool to model the current and future potential geographic distribution of *Kalopanax septemlobus* (Araliaceae) in South Korea. We then applied a graph-theoretical method to predict the likely impacts of climate change on habitat connectivity. Natural populations of *K. septemlobus* are threatened by climate change and from illegal logging for timber, medicine, and edible products (Chang et al., 2001; Kang, 2003). The objectives of this study are: (1) to identify climatic and topographic factors associated with *K. septemlobus* distribution by using ecological niche modeling (ENM), (2) to predict the current distribution of suitable habitats and project them under future climate scenarios for 2050, and (3) to evaluate changes in habitat distribution patterns and connectivity.

## 2. Materials and methods

### 2.1. Study area and focal species

The study area is in the southern half of the Korean Peninsula and the islands of South Korea (Fig. 1), which lie between latitudes 33° and 39°N, and longitudes 124° and 131°E. Its total area is 100,148 km², approximately 64% of which is covered by forests mostly in the north and east regions (KFS, 2012). The area is in a temperate zone with four distinct seasons and is affected by East Asian monsoons. However, global warming has increased the temperature and precipitation levels and widened seasonal and regional weather differences on the Korean Peninsula, changing its climate gradually closer to a subtropical climate (Philander, 2012).

*K. septemlobus*, commonly known as the prickly castor oil tree, is a hermaphroditic, deciduous tree species in the family Araliaceae. Widely but sparsely distributed through Northeast Asia (Lee and Kang, 2002; Ohashi, 1994), this species blooms in July–August, and various insects act as agents of pollination (Fujimori et al., 2006). Fruits are available in September and October, and seeds are dispersed by birds and squirrels (Iida and Nakashizuka, 1998). It is a multi-purpose tree, important for high quality timber and as a source of food and medicine. However, illegal cutting and over-exploitation due to the increasing demand have led to damage and destruction of its natural habitat (Kang, 2003).
2.2. Habitat distribution modeling under current and potential future climates

The present potential distribution of *K. septemlobus* was constructed by relating occurrence locations to bio-climatic and topographic variables through a presence-only machine-learning maximum entropy (MaxEnt) distribution modeling (Phillips et al., 2006). We obtained 72 occurrence records of *K. septemlobus* from previously published studies (Kang, 2003; Lee et al., 2000; ME, 2002; Sakaguchi et al., 2012a, 2012b) (Fig. 1). The occurrence locations were separated by a minimum distance of 1 km to avoid spatial autocorrelation biases (Legendre et al., 2002).

Eight environmental variables were chosen according to their potential relevance to the *K. septemlobus* habitat distribution based on previous research (Sakaguchi et al., 2012a, 2010) and other habitat modeling studies (e.g., Khanum et al., 2013; Kumar and Stohlgren, 2009). Four bioclimatic variables, including annual mean temperature, mean diurnal temperature range, precipitation of warmest quarter, and precipitation of coldest quarter, at a 30 arcsecond (ca. 1 km) spatial resolution were obtained from the WorldClim database for the current climate of 1950–2000 (Hijmans et al., 2005; http://www.worldclim.org/bioclim.htm). For topographic variables, the images in the study area, the bioclimatic variables were resampled to a 100 m resolution using bilinear interpolation. We also used four topographic variables including slope and aspect, both in degrees; solar radiation; and topographic position index derived from a 100 m digital elevation model. Potential solar radiation was calculated as the sum of direct, diffuse, and reflected radiation for one year based on terrain shading (Kumar et al., 1997). The topographic position index was generated by using ArcView 3.3 with the Topographic Position Index extension version 1.3a (Jenness, 2006). MaxEnt determines the relative contribution of each variable to the model (Phillips et al., 2006). Only variables contributing 1% or more to the model were included in the final model. We tested for multicollinearity by examining Spearman’s rank cross-correlations among all of the variables based on 10,000 randomly generated points within the study area. No variables were strongly correlated (i.e., $r_s < 0.6$).

We performed MaxEnt modeling by using the default settings, with 1000 iterations. Thirty model replicates were processed; 70% of the locations were randomly selected each time to train the model, and the remaining 30% were used to test the model’s predictions. We used ENMTools 1.3 (Warren et al., 2010; Warren and Seifert, 2011) to select the most parsimonious model based on the corrected Akaike’s information criterion (AICc) scores (Burnham and Anderson, 2002). Using the 10 percentile training presence logistic threshold ($>0.295$) (Escalante et al., 2013) from the best-fitting model, habitat patches were defined as contiguous pixels based on an eight-neighbor rule. Patches less than 5 ha, the best-fit prediction of organisms) within a landscape-ecological context (Minor et al., 2009). A graph is a set of nodes (i.e., discrete patches) connected by links (i.e., movement of organisms) within a landscape-ecological context (Minor and Urban, 2008).

Habitat networks were analyzed as undirected complete graphs. Link weights were determined on the basis of Euclidean edge-to-edge distances between habitat patches. Although long-distance seed dispersal events for *K. septemlobus* are rare (N. Fujimori, unpublished data), we used different threshold distances (100 m–25 km) to consider a range of possible pollen and seed dispersal distances.

To compare current and future levels of overall habitat connectivity, we estimated the Probability of Connectivity (PC) (Saura and Pascual-Hortal, 2007), which is computed as:

For each of the three habitat models, including the current *K. septemlobus* distribution and the two predictions under the RCP scenarios, we calculated the total area of suitable habitats. Then, we estimated the amounts of stable, unsuitable, lost, and gained habitats for each of the two future models. Stable habitat refers to the areas of the current potential range predicted to remain suitable in 2050, and lost habitat refers to areas not predicted to remain suitable in the same period. Gained habitat includes areas that are predicted to be suitable in 2050 that are not currently suitable.

To measure habitat fragmentation, we computed the number of patches (NP), area weighted mean patch size (AWMPS), and largest patch index (LPI), using FRAGSTATS (version 4.2; McGarigal et al., 2012). The LPI is the percentage of total landscape area comprised by the largest patch. The AWMPS is the sum of patch areas across all patches multiplied by the proportional abundance of the patch (i.e., patch area divided by the sum of patch areas). AWMPS is considered more robust than the simple mean patch size when quantifying changes in landscape structure over time (Li and Archer, 1997). We also analyzed altitudinal range shifts of the species by using the mean elevation of current and future suitable habitats.

We estimated the degree of overall connectivity by using a graph-theoretical approach. In ecology, connectivity is often analyzed by graph-theoretical methods with minimal data inputs, which show promise for providing functional and ecologically relevant measures of landscape structure and process (Kang et al., 2012; Minor and Urban, 2007; Urban et al., 2009). A graph is a set of nodes (i.e., discrete patches) connected by links (i.e., movement of organisms) within a landscape-ecological context (Minor and Urban, 2008).
where $n$ is the total number of habitat patches, $a_i$ and $a_j$ are the areas of patches $i$ and $j$, respectively; and $A_L$ is the total landscape area. $p_{ij}^*$ is defined as the maximum product probability of all possible paths between the patches $i$ and $j$. The product probability of a path is the product of all the $p_{ij}$ (link weights) included in the path. The $p_{ij}$ values (dispersal probability) were calculated by using a negative exponential function of inter-patch distance (Bunn et al., 2000; Urban and Keitt, 2001), with $p_{ij} = 0.05$ for a threshold distance. When $i = j$, the $p_{ij}^*$ equals 1. PC is defined as the probability that two organisms randomly placed in a landscape will be interconnected by falling into habitat areas within reach of each other given a set of habitat nodes and links between them (Saura and Pascual-Hortal, 2007). The PC index may be an appropriate metric for studying overall flows of organisms irrespectively of their origin (Bodin and Saura, 2010).

By comparing the PC values of the current and future habitat networks, we predicted the potential impact of climate change on the connectivity of K. septemlobus habitats. Based on the present PC values under current climate conditions, we calculated the percent of decrease in PC connectivity caused by shifts in suitable habitat range under RCPs 4.5 and 8.5 for 2050. Graphab 1.2 was used for constructing and evaluating the habitat networks (Foltête et al., 2012).

3. Results

The best model for predicting the K. septemlobus occurrence probability had a good fit (AUC-train = 0.89; AUC-test = 0.85). Among the eight variables, the most influential predictor was average annual mean temperature (65.3%); (2) slope (16.8%); (3) precipitation of warmest quarter (5.8%); (4) precipitation of coldest quarter (4.5%); (5) mean diurnal temperature range (2.4%); (6) aspect (2.3%); (7) solar radiation (1.6%); and (8) topographic position index (1.1%). The occurrence probability of K. septemlobus decreased with increasing annual mean temperature (Fig. 2a), while it generally increased with increasing slope (Fig. 2b).

The current suitable habitats, covering 14.2% of the total land area, were mainly found in the protected regions of the Baekdu-daegan, the eastern part of Gyeongbuk province, and the west and central regions of Jeju Island (Fig. 3a). However, more than 50% of current habitat was expected to be rendered unsuitable under RCPs 4.5 and 8.5 (Fig. 3b, c). Particularly in the central and southern regions of South Korea, most patches of suitable habitat were projected to disappear under future climatic conditions (Fig. 3b, c). Under an intermediate scenario, we detected a habitat loss of 57.6% and a habitat gain of 6.3%, resulting in net loss of 51.3% (Table 1). Under a high scenario, we detected a habitat loss of 85.9% and a habitat gain of 0.9%, resulting in net loss of 85.0% (Table 1). Those scenarios also decreased the number of suitable patches in RCPs 4.5 and 8.5 by 51.4% and 83.4%, the size of the largest patch by 62.3% and 96.1%, and the area weighted mean patch size by 69.7% and 97.4%, respectively (Table 2). The mean elevation of future suitable habitats was increased by 88 m and 162 m in RCP 4.5 and RCP 8.5, respectively, from the current mean elevation of 648 m.

The connectivity of the predicted future networks was significantly lower than that of the current network at all threshold distances. In addition, the decreases in connectivity under RCP 8.5 were much higher than that under RCP 4.5 (Fig. 4); more than 99% reduction in connectivity was detected under RCP 8.5 at all threshold distances. Under RCP 4.5 for 2050, the current connectivity level was reduced by almost 77% for the largest dispersal distances and even more for shorter dispersal distances. The average values for the percentage decrease in connectivity over all distances were 78.4% and 99.4% in 2050 under RCPs 4.5 and 8.5, respectively. Moreover, the percentage decreases in network connectivity were consistently higher than decreases in suitable habitat area (i.e., 51.3% and 85.0% for RCPs 4.5 and 8.5, respectively). This result suggests that the loss in habitat area had more deleterious effects on network connectivity.

4. Discussion and conclusions

In this study, we used species distribution modeling coupled with habitat pattern and network analyses to quantify changes in the extent and connectivity of the K. septemlobus habitat under current and projected climate conditions for 2050. We found that the distribution of K. septemlobus is primarily determined by average annual temperature, indicating that increasing average temperatures due to climate change can be particularly detrimental to the species. As a result, future species distribution models showed that its geographic distribution in South Korea would

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**Fig. 2.** Response curves of Kalopanax septemlobus to gradients of (a) annual mean temperature and (b) slope. Each of the curves represents a MaxEnt model created using only the corresponding variable.
shrink under predicted levels of climate warming. In addition, the resulting changes in the amount of available suitable habitat had a large impact on overall connectivity. If warming continues unabated, most populations of *K. septemlobus* in South Korea will become physically isolated from one another. Therefore, for this species, a rapid rise of mean temperature will become a severe stress factor. The effects will be particularly extreme if the species lacks capacity to move to other suitable habitat areas and if a lack of gene flow among fragmented populations occurs with the lessened capacity to adapt (Angert et al., 2011; Schloss et al., 2012).

**Table 1**  
Current and predicted suitable habitats for *Kalopanax septemlobus* based on two Representative Concentration Pathway (RCP) scenarios, RCPs 4.5 and 8.5, for the year 2050.

<table>
<thead>
<tr>
<th>Scenario</th>
<th>Total Habitat Area (km²)</th>
<th>Stable Habitat Area (km²)</th>
<th>Lost Habitat Area (km²)</th>
<th>Gained Habitat Area (km²)</th>
<th>Net Habitat Loss Area (km²)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Current</td>
<td>14,299.4</td>
<td>-</td>
<td>-</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>RCP 4.5</td>
<td>6961.7</td>
<td>6059.5</td>
<td>8239.9</td>
<td>902.2</td>
<td>7317.7</td>
</tr>
<tr>
<td>RCP 8.5</td>
<td>2139.2</td>
<td>12,160.2</td>
<td>0.9</td>
<td>12,060.2</td>
<td>85.0</td>
</tr>
</tbody>
</table>

* Percentages were obtained against the current potential range.

**Table 2**  
Fragmentation metrics for the current and future suitable habitats for *Kalopanax septemlobus* based on two Representative Concentration Pathway (RCP) scenarios, RCPs 4.5 and 8.5, for the year 2050.

<table>
<thead>
<tr>
<th>Scenario</th>
<th>Number of Patches</th>
<th>Area Weighted Mean Patch Size (km²)</th>
<th>Largest Patch Index</th>
</tr>
</thead>
<tbody>
<tr>
<td>Current</td>
<td>10,671</td>
<td>4247.8</td>
<td>7.7</td>
</tr>
<tr>
<td>RCP 4.5</td>
<td>5187</td>
<td>1288.1</td>
<td>2.9</td>
</tr>
<tr>
<td>RCP 8.5</td>
<td>1760</td>
<td>108.9</td>
<td>0.3</td>
</tr>
</tbody>
</table>
The MaxEnt model showed that annual mean temperature and slope were the two most important variables predicting habitat suitability for *K. septemlobus*, but variation in the annual mean temperatures explained more than 65% of the total variation in its distribution. This concurs with the results of a previous study suggesting that the ecological niche of this species is constrained mainly by a temperature component among the bioclimatic variables (e.g., Sakaguchi et al., 2010). In regard to slope, some studies reported that *Kalopanax* stands were located in relatively steep and ridge-slope areas (e.g., Kang and Lee, 1998; Lee et al., 2000), possibly as a result of interspecific competition. Although the growth rate of *K. septemlobus* is high in low-slope valley areas where soil moisture and nutrient are rich, it is known that the species is usually overcome by other competitive tree species and is released to ridge-slope areas (Lee and Kang, 2002).

The predicted current species distribution supported an earlier finding that *K. septemlobus* is widespread both horizontally and vertically in temperate deciduous or coniferous forests of South Korea (Fig. 3a) (Lee and Kang, 2002; Sakaguchi et al., 2012a). However, the model’s predictions indicated that this species had a very low probability (<0.25) of occurring in areas with a moderate to high mean annual temperature of 10°C–16°C (Fig. 2a). Thus, the species may become increasingly vulnerable to thermal stress associated with climate warming (Allen et al., 2010). Moreover, the model results indicated that persistence of the species may largely depend on its ability to adapt locally to such pressure, which needs further exploration.

With an increase in temperature, the potential range of the species under present climatic conditions would become unsuitable and would shift northwards or to higher elevations (Sakaguchi et al., 2010). The estimated rates of elevational range shift in suitable habitats under the RCPs 4.5 and 8.5 for 2050 as a function of threshold distance (i.e., dispersal ability).

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In the case of short distance dispersers such as *K. septemlobus*, the isolation of a potentially suitable habitat located at the center of a species’ geographic range raises a concern on the effectiveness of habitat networks because the ability of the species to disperse and colonize newly suitable areas would be restricted (Mazaris et al., 2013). Although climate warming will greatly affect populations inhabiting the southern edge of their distribution ranges (Jueterbock et al., 2014), it can also further weaken the capacity of habitat networks to sustain metapopulations of species with short-range dispersal. For these species, large clusters of functionally connected patches would be broken into smaller and more isolated suitable habitats within and surrounding the central parts of the habitat network, currently acting as stepping-stone corridors connecting the large clusters, become smaller and more isolated in response to climate warming. In this sense, the results reported in this study, i.e., the highest reduction in habitat connectivity at lower dispersal distances (Fig. 4), suggest that species with short-range dispersal are more sensitive to habitat connectivity loss under climate change than species with long-range dispersal.

*K. septemlobus* is a long-lived perennial tree. Long-lived species may persist in fragmented landscapes for long periods during which opportunities for long-distance dispersal of either seed or pollen would increase. Long-lived species also have the potential for rapid adaptation if they have high genetic diversity and large population sizes, whereas short-lived species can evolve rapidly because of their quick generation replacement times. Typical populations of *K. septemlobus* in Korea are small and are known to have very narrow genetic diversity (Sun et al., 2012). Thus, adapting to future climate conditions will be difficult for *K. septemlobus*, particularly if the rate of warming is too rapid. Moreover, considering the dispersal capabilities of *K. septemlobus*, this species will likely be unable to track the projected changes in climate over this century (Sakaguchi et al., 2010).

The complete disappearance of *K. septemlobus* is hardly anticipated, even in areas with decreased habitat suitability by 2050, when considering both the longevity of the species and the competition with other species for limited space (Sakaguchi et al., 2010). For several plant species, climatic responses often lag behind climate change (Bertrand et al., 2011). Although the estimates provided here should not be taken as precise predictions, climate-induced range contraction of *K. septemlobus* is expected to occur throughout the country within the next few decades.

Our results may be conservative because we did not account for the effects of land-use change and elevated CO₂ on *K. septemlobus* habitats. Climate change and agriculture are interrelated processes that threaten biodiversity (MA, 2005). Temperature increases will lead agricultural lands suitable for crops to expand into higher elevational areas. This conversion of natural vegetation to agriculture at high elevations will be an important source of CO₂ emissions. Although elevated CO₂ levels alone may benefit the growth of *K. septemlobus* individuals (Watanahe et al., 2010), *K. septemlobus* will likely experience increases in acute heat stress under elevated CO₂, which can cause large-scale forest mortality events and overall habitat connectivity. Furthermore, the percentage of connectivity loss for each scenario was higher than that expected from the variation in the available suitable habitat area alone or in each habitat pattern metric. This indicates a significant decrease in structural connectivity following habitat loss and fragmentation. Thus, continued warming may create small, isolated populations much more sparsely distributed than those under present conditions. Because most seeds of the species are dispersed less than 1 km (N. Fujimori, unpublished), such populations should undergo very limited seed dispersal. In turn, they may be more susceptible to demographic and genetic stochasticity (Gilpin and Soulé, 1986; Goodman, 1987; Shaffer, 1981).
subsequent decreases in ecosystem productivity and biodiversity (Ciais et al., 2005; Hamilton III et al., 2008; Thomas et al., 2004). For more accurate future predictions of species distributions, the impacts of the interacting pressures of land use change, elevated CO2, and climate warming on species should be explored.

K. septemlobus is of ecological and medical importance in Korea. However, over-exploitation combined with climate change will likely lead to its overall population decline, as discussed by Midgley et al. (2003). In addition, it is likely that its remaining range will become disjunct and future metapopulations will become isolated. If a species is not phenotypically plastic, it will be confronted with the high pressure of adapting through evolution or moving through range shifts. And, if a species is unable to do either of these quickly or well, future extinction is likely. It is known that evolutionary responses may be insufficient to keep pace with anthropogenic climate change (Franks et al., 2014). Hence, identifying “climate change refugia”, areas that are suitable under both current and future climates (Franklin et al., 2013), in addition to maintaining or improving connections between the areas to enable range shifts, is important for ensuring species’ long-term persistence. For areas at high risk of habitat loss that are fairly isolated from climate change refugia (i.e., mountain ranges in the southern part of the study area), adaptive strategies such as assisted migration may be implemented (Vitt et al., 2010). Finally, the predictive modeling approach used in this study may also be useful to assess the vulnerability of other threatened species and their habitats to climate change.

Author contributions

W.K. and D.L. designed the study; W.K. and C.-R.P. collected the data; W.K., E.S.M., and C.-R.P. analyzed the data; and W.K. wrote the paper, with substantial contributions from all authors. All authors have read and approved the final manuscript.

Acknowledgments

This study was supported by the Basic Science Research Program through the National Research Foundation of Korea (NRF) funded by the Ministry of Education (No. 2011-0024289).

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