# ORIGINAL RESEARCH

# WILEY Ecology and Evolution

# Host plant density and patch isolation drive occupancy and abundance at a butterfly's northern range margin

Yoan Fourcade 💿 📔 Erik Öckinger 💿

Department of Ecology, Swedish University of Agricultural Sciences, Uppsala, Sweden

#### Correspondence

Yoan Fourcade, Department of Ecology, Swedish University of Agricultural Sciences, Uppsala, Sweden. Email: yoanfourcade@gmail.com

#### Funding information

Swedish Research Council, Grant/Award Number: 621-2010-5589; Signhild Engkvist Foundation; Lunds Djurskyddsfond; Royal Swedish Academy of Agriculture and Forestry; Department of Ecology SLU

#### Abstract

Marginal populations are usually small, fragmented, and vulnerable to extinction, which makes them particularly interesting from a conservation point of view. They are also the starting point of range shifts that result from climate change, through a process involving colonization of newly suitable sites at the cool margin of species distributions. Hence, understanding the processes that drive demography and distribution at high-latitude populations is essential to forecast the response of species to global changes. We investigated the relative importance of solar irradiance (as a proxy for microclimate), habitat quality, and connectivity on occupancy, abundance, and population stability at the northern range margin of the Oberthür's grizzled skipper butterfly Pyrgus armoricanus. For this purpose, butterfly abundance was surveyed in a habitat network consisting of 50 habitat patches over 12 years. We found that occupancy and abundance (average and variability) were mostly influenced by the density of host plants and the spatial isolation of patches, while solar irradiance and grazing frequency had only an effect on patch occupancy. Knowing that the distribution of host plants extends further north, we hypothesize that the actual variable limiting the northern distribution of P. armoricanus might be its dispersal capacity that prevents it from reaching more northern habitat patches. The persistence of this metapopulation in the face of global changes will thus be fundamentally linked to the maintenance of an efficient network of habitats.

#### KEYWORDS

climate change, habitat quality, land-use, metapopulation, microclimate, peripheral populations

# 1 | INTRODUCTION

Populations located at the periphery of species' ranges have a particular value for conservation (Lesica & Allendorf, 1995). The abundance of a species typically decreases toward the edge of its range (Brown, 1984; Brussard, 1984; but see Sagarin & Gaines, 2002). Also, peripheral populations can typically only occupy a fraction of the potential habitat due to climatic constraints, resulting in more fragmented populations at the periphery than at the core of the range (Thomas, 1993). In combination, this makes marginal populations more prone to extinction (Hardie & Hutchings, 2010). In addition, populations living at the latitudinal margins of a species' range are critical in the process of species response to climate change. Leading-edge populations, being located at the potential colonization front, are those on which the capacity for a species to shift or expand its distribution relies (Hampe & Petit, 2005; Thuiller et al., 2008). Providing that all other requirements are met, range expansion can occur by recurrent poleward dispersal events from these populations followed by population growth in newly colonized sites (Hampe & Petit, 2005; Smale & Wernberg, 2013). Therefore, identifying the drivers of dynamics of

This is an open access article under the terms of the Creative Commons Attribution License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited.

 $\ensuremath{\mathbb{C}}$  2016 The Authors. Ecology and Evolution published by John Wiley & Sons Ltd.

WILEY\_Ecology and Evolution

FOURCADE AND ÖCKINGER

high-latitude populations is essential to understand the factors that shape range limits and to forecast the response of species to climate change. Ultimately, this knowledge is crucial to inform the conservation of these vulnerable populations.

Beyond climate, several biotic and abiotic factors have the potential to drive population processes at latitudinal range margins and to determine the limits of species' ranges (Sexton, McIntyre, Angert, & Rice, 2009). For example, although there is ample evidence that many species are currently responding to climate change by shifting their distribution poleward (Chen, Hill, Ohlemuller, Roy, & Thomas, 2011; Gillings, Balmer, & Fuller, 2015: Hickling, Roy, Hill, Fox, & Thomas, 2006: Parmesan et al., 1999), the actual patterns of range shifts have been shown to result from a complex interaction between climate, biotic interactions (Van der Putten, Macel, & Visser, 2010), intrinsic species traits, and anthropogenic pressures (Jetz, Wilcove, & Dobson, 2007). In this regard, habitat fragmentation caused by human land use can be a key limiting factor. It may significantly impact range shift opportunities (Burrows et al., 2014) and can accelerate the extinction of isolated populations by disconnecting them from other suitable areas (Opdam & Wascher, 2004). Local habitat quality can also interact with climate variables in determining range boundaries and the response of populations to climate change (Kleijn et al., 2010; Nicolè, Dahlgren, Vivat, Till-Bottraud, & Ehrlén, 2011; Seabrook et al., 2014). Biotic interactors (prey, predators, or hosts) are important determinants of habitat quality and can strongly influence population performance and species distribution (Louthan, Doak, & Angert, 2015). They can be so important that a mismatched response to climate change can limit range shifts that would have otherwise occurred if species tracked solely their climate niche (Pelini et al., 2009; Schweiger, Settele, Kudrna, Klotz, & Kuhn, 2008).

One approach to determine the relative importance of these factors is to study their impact on the dynamics of high-latitude populations. In this regard, population responses to different microclimates offer an indirect but useful assessment of the climate-related reaction norm of the species (Lawson, Bennie, Hodgson, Thomas, & Wilson, 2014; Thomas, 1993). If microclimate is the main driver of the abundance or distribution of these marginal populations, it would suggest that the species' range is likely limited by climatic factors. We can thus expect the species to react to climate change by shifting its range poleward (Bennie et al., 2013). If, instead, habitat fragmentation is already limiting current patterns, restoring landscape connectivity may be pivotal to the conservation of these populations. Otherwise, it is unlikely that the species will be able to cope with climate change and expand its range unless new habitats are created. Similarly, if population demography and distribution at leading-edge margins closely depend on such biotic factors, the response of the focal species to future changes will largely be driven by the response of its co-occurring species (Gilman, Urban, Tewksbury, Gilchrist, & Holt, 2010; Van der Putten et al., 2010). Whether these populations can be the starting point of climate change tracking thus depends on their fine-scale drivers of distribution and dynamics and on the species intrinsic habitat requirements.

Here, we examined the effect of variation in solar irradiance (used as a proxy for potential microclimate), patch quality, or connectivity across a network of patches at a butterfly's northern range margin. We



**FIGURE 1** Adult Oberthür's grizzled skipper (*Pyrgus armoricanus*). Photograph by Theresia Widhalm and Alexander Neubauer

used a habitat network including a large part of the total population of Oberthür's grizzled skipper (Pyrgus armoricanus) (Figure 1) in Sweden as a model to infer the importance of microclimate, patch quality, and connectivity on the regional distribution and local abundance at a species' northern range margin. The dynamics of P. armoricanus in southern Sweden has been proposed to be driven by metapopulation processes (Öckinger, 2006). Following metapopulation theory (Hanski, 1998), we expect the spatial configuration of patches-their area and their degree of isolation from the surrounding patches-to be an important driver of P. armoricanus abundance and habitat occupancy. This would highlight the importance of dispersal opportunities and thus the decisive impact of human land use on the current conservation status of the species and on its future response to climate change. Similarly, as vegetation structure is known to affect butterflies in general (Kruess & Tscharntke, 2002), and this species in particular (Eilers, Pettersson, & Öckinger, 2013), we also investigated the effect of variable grazing intensities on interpatches variation in occupancy and abundance. Moreover, we know from a previous study that microclimate influences the choice of oviposition sites in this species (Eilers et al., 2013). This factor is thus likely to affect the observed probability of habitat patches to be occupied and the population size they can sustain. Finally, owing to the fact that the presence of P. armoricanus in a grassland patch is generally closely linked to the availability of its host plant species (Öckinger, 2006), we also assessed the effect of the density of hosts. By ranking the importance of each of these factors in explaining variation in occupancy and abundance among habitat patches, we can gain insights into the population processes acting at high-latitude range margins and more specifically the potential response of this species to climate change.

## 2 | METHODS

#### 2.1 | Study area and data collection

Our study species, Oberthür's grizzled skipper (*P. armoricanus*) (Figure 1), has a wide but fragmented distribution throughout North

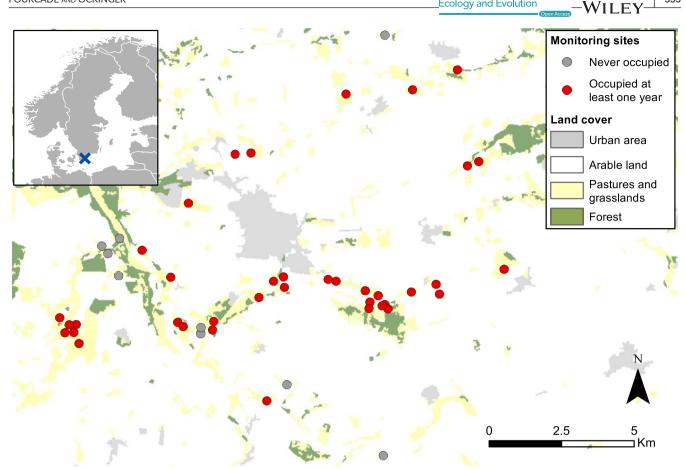


FIGURE 2 Location of the study area (shown as a cross in upper left inset) and spatial configuration of the 50 sites surveyed in this study

Africa and Europe. Its northernmost populations are located in southern Scandinavia (Sweden and Denmark), in a relative isolation from other populations in western and central Europe (Kudrna et al., 2011). Its habitat consists of seminatural grasslands which host the specific plant species where it lays its eggs and on which its larvae feed. Scandinavian populations are known to select primarily Filipendula vulgaris and Helianthemum nummularium (Christensen, 2000; Eilers et al., 2013), two species that are known to occur in a fragmented distribution up to 600 km further north (Hultén, 1971). The species has two generations per year: Spring generation adults fly from mid- or late May to mid-June, and the summer generation flies in August. In addition to Denmark, its Scandinavian distribution is restricted to a small area of ca. 30 × 20 km in southern Sweden where it occurs in a network of small and fragmented patches (Öckinger, 2007). There exist no records of historical occurrences further north (Eliasson, Ryrholm, Gärdenfors, Holmer, & Jilg, 2005; Nordström, Opheim, & Valle, 1955). The habitat patches analyzed in this study (min area: 0.028 ha, max area: 14.71 ha), defined as patches of dry unfertilized grasslands with the presence of at least one of the host plants F. vulgaris and H. nummularium (Öckinger, 2006), were located in the core of this system, mainly around the town of Tomelilla (Figure 2). Adjacent habitat patches were defined as discrete if separated by at least 50 m of divergent vegetation (often arable fields or agriculturally improved grassland) as recommended by Ojanen, Nieminen, Meyke, Poyry, and Hanski (2013).

The occurrence and abundance of P. armoricanus were monitored in 50 habitat patches from 2004 to 2015. Between 2004 and 2011, sites were surveyed twice a year to record the abundance of both generations. From 2012, only the summer generation was surveyed. Of the 50 patches, 34 were surveyed all 12 years, eight sites 11 years, six during 9 years, one during 8 years, and one site during 7 years only. Sites were monitored by slowly walking a transect (10 m width) covering the entire area of the patch. As transects were designed to allow the entire patch area to be surveyed, the transect length was proportional to patch area, but, the sampling effort per unit area was constant. Counts should thus reflect butterfly abundance within patches and not only transect length. All observed adult P. armoricanus individuals were recorded. If necessary, species identity was confirmed by capturing individuals with a handheld net. As we observed that the abundance of the summer generation is on average almost three times higher than the spring generation (Table 1), we considered that processes acting in each of them might be different (see, for example, Roy & Thomas, 2003) and thus analyzed each generation separately. Therefore, we derived for each habitat patch and each generation: occupancy and the average and variability of abundance. Occupancy was expressed, separately for spring and summer generations, as the number of years a patch was occupied divided by the number of years it was surveyed in this generation. As imperfect detection can bias estimates of butterfly abundance and occupancy, it is sometimes advised to use multiple surveys per season to accurately estimate occupancy rates (MacKenzie WILFY\_Ecology and Evolution

**TABLE 1** Average characteristics of the 50 sites surveyed. Response variables: mean (±*SD*) occupancy, average, and coefficient of variation (CV) of abundance per patch for each generation. Explanatory variables: mean (±*SD*) connectivity (depends on abundance of other patches, hence one value for each generation), area, host plant density, mean and standard deviation of solar irradiance per site, and number of sites in each category of grazing

	Spring generation	Summer generation		
Dependent variables				
Occupancy	0.47 (±0.39)	0.52 (±0.40)		
Average abundance <sup>a</sup>	6.13 (±10.39)	16.16 (±27.83)		
CV abundance <sup>a</sup>	1.53 (±0.59)	1.62 (±0.73)		
Explanatory variables				
Connectivity	4.46 (±7.87)	11.80 (±19.41)		
Area (m <sup>2</sup> )	15,756 (±24,882)			
Host plant density (%)	3.61 (±5.63)			
Grazing				
Never	7			
Sometimes	19			
Always	24			
Mean solar irradiance (Wh/m²)	807,542 (±48,325)			
SD solar irradiance (Wh/m²)	47,678 (±31,754)			

<sup>a</sup>Excluding sites that were never occupied during the 12 years of survey.

et al., 2002). Instead, we chose here, due to time limitations, to maximize the number of patches visited each year rather than visiting each patch multiple times. However, we accounted for possible incomplete detection by considering a patch unoccupied only when no individuals were recorded during two consecutive surveys, whether these surveys occurred in the same year (spring and summer generations) or in different years (summer generation at year t and spring or summer generation-when only summer generations were surveyed-at year t + 1). This conservative estimate should ensure the robustness of our results even if some sites were mistakenly assumed to be unoccupied. Abundance was measured only for patches that have been recorded as occupied in at least one survey, regardless of the generation. This means that, for example, if a patch was only occupied in one summer generation survey, it was still retained for the abundance analyses of the spring generations, where it thus had a mean abundance of zero. Abundance variability was defined as the coefficient of variation of abundances across years.

We characterized each habitat patch by six variables that described habitat quality, solar irradiance, or spatial configuration of patches (Table 1). In each habitat patch, we estimated the density of *F. vulgaris* and *H. nummularium*. This was done by randomly placing 10 quadratic  $1 \text{ m}^2$  plots along the butterfly monitoring transect, and estimating the percentage cover of each host plant separately within each plot. Densities of host plants and flowers were recorded in May-June 2010. As both host plants are perennial and their densities have

remained relatively stable over time (E. Öckinger, personal observation), the records from 2010 are assumed to represent the entire study period. P. armoricanus females show similar preferences for F. vulgaris and H. nummularium, although the former plant species is typically much more abundant (Eilers et al., 2013). Therefore, we pooled the cover of F. vulgaris and H. nummularium and used the averaged value over the 10 plots as a measure of host plant density per patch. All patches had a density of host plants between 0% and 15%, except one habitat patch that showed an exceptionally high host plant density of ca. 34%. We also categorized each patch by its grazing frequency according to three classes: sites that were never grazed during the whole period of survey (thereafter referred to as "never"), sites that were grazed every year ("always") and sites that were grazed or not depending on the year ("sometimes"). We used solar irradiance to characterize potential microclimate, a feature that is known to affect P. armoricanus oviposition site selection (Eilers et al., 2013). Solar irradiance depends on latitude, elevation, aspect, slope, and surrounding topography and reflects the level of energy that is received at a given point of Earth. We used a digital elevation model from the Swedish Lantmäteriet (2015), produced by laser scanning. The accuracy of the elevation model is 0.5 m, and the resolution of grid cells is 2 m. Insolation was estimated as the total direct solar irradiance per 2 m grid cell per year ( $Wh/m^2$ ), using the solar radiation function in the Spatial Analyst toolbox in ArcGIS 10.2 (ESRI Inc., Redlands, CA, USA), based on latitude, slope, aspect, and effects of shading from the elevations of surrounding cells. We calculated the mean and standard deviation of solar irradiance for each habitat patch to represent the average and variability in microclimate.

We also quantified for each patch the two main predictors of metapopulation dynamics: area and connectivity. The boundaries of habitat patches were defined in the field and digitized in a GIS. To account for the fact that transect lengths were dependent on patch area (see monitoring protocol above), we included area as a covariable in all analyses. Patch areas were calculated using ArcGIS 10.2. As a measure of connectivity, we used the index S; developed by Hanski (1999) and calculated as  $S_i = \sum_{i \neq i} e^{-\alpha d_{ij}} N_i$ .  $S_i$  estimates the connectivity at patch *i*, where  $d_{ii}$  is the Euclidian distance between patches *i* and *j* (in meters) and  $N_i$ the observed abundance at patch *j* (reflecting the number of potential emigrants from patch j),  $\alpha$  being a constant describing how fast immigration probability from patch *j* decreases with increasing distance. Here,  $\alpha$  was set as 0.0034, corresponding to an average movement distance of 295 m, according to a previous analysis of mark-recapture data (E. Öckinger, unpublished data). When calculating connectivity, we took into account not only the 41 patches analyzed in this study, but all potential habitat patches within a 35 × 35 km square including the entire known distribution of P. armoricanus in Sweden (25 patches in addition to the 50 surveyed patches). As this index depends on abundance of all surrounding patches, we calculated S<sub>i</sub> for each year and generation, and used in further analyses its average value over all years for each generation separately. When abundance data were unavailable in a given year, we used the average abundance at this site across all years of survey. Similarly, for all sites that were not part of the annual monitoring, we used a rough estimation of abundance

WILEY-

at these sites based on a systematic mapping of the species' entire distribution in Sweden in 2007 and 2010.

#### 2.2 | Statistical analyses

We analyzed the effect of habitat quality, microclimate, area, and connectivity on patch occupancy and the average and variability of butterfly abundance in both generations separately. Occupancy was analyzed by a generalized linear model with binomial error distribution and logit link. The response variable was defined, for each generation, as the number of years a patch was occupied divided by the number of years it was surveyed. To account for potentially false absences, we took a conservative approach and only considered a patch as unoccupied if no P. armoricanus individuals were observed there during two consecutive surveyed generations; otherwise, the patch was considered as being still occupied. We used linear regressions to model average and variability of abundance per site. The average and coefficient of variability of abundance were log-transformed prior to analyses so that model predictions fall in the interval  $[0, +\infty]$ . Moreover, a small value (0.1) was added to all abundance data so that the sites that were never occupied could be included as well (log-transformation impossible when average abundance = 0). As explanatory predictors, we used two variables linked to the spatial configuration of sites: patch area and connectivity, two variables describing habitat quality: host plant density and grazing frequency (defined as three categories: always, never, or sometimes grazed during the survey period), and two variables describing microclimate: mean and standard deviation of solar irradiance. In addition, we aimed to test whether the proximity of neighboring patches, and thus a higher immigration probability in a metapopulation model, could balance low habitat guality or unfavorable microclimatic conditions. Therefore, we also included in all models as explanatory variables the two-way interactions between connectivity and site quality variables (host plant density and grazing) or microclimate (mean and SD of solar irradiance), resulting in four additional predictors.

We adopted an information-theoretic approach (Burnham & Anderson, 2002) by computing models with all combinations of variables, and ranked them by their second-order Akaike information criterion (AIC<sub>c</sub>). A multimodel inference was then performed by averaging all models whose cumulative Akaike weight was <0.95 (Burnham & Anderson, 2002). We extracted standardized averaged parameter estimates of all variables and interactions and estimated relative variable importance based on the sum of Akaike weights of all candidate models containing the variable. Multimodel inferences were run using the "MuMIn" package (Barton, 2013) in R 3.2.2 (R Development Core Team, 2015). Partial relationships were visualized by plotting model predictions against variations in the variable of interest while holding all other variables at their median-grazing frequency being set to "sometimes." For visualizing interactions with connectivity, this approach was performed for three levels of connectivity corresponding to its 0.25, 0.5, and 0.75 quantiles.

# 3 | RESULTS

Nine patches were occupied through all 12 years, nine patches were never occupied, and 32 patches were occupied during at least 1 year. Patch occupancy thus ranged from 0 to 1 (mean =  $0.51 \pm 0.39$  SD) when both generations where considered together, with roughly similar values in the spring (from 0 to 1, mean =  $0.47 \pm 0.39$  SD) and the summer generations (from 0 to 1, mean =  $0.52 \pm 0.40$  SD) (Appendix 1 and Table 1). The average abundance per patch (both generations: from 0.053 to 98.05. mean = 12.50 ± 21.34 SD) was generally higher during the summer generation (from 0 to 128.75; mean =  $16.16 \pm 27.83$  SD) than during the spring generation (from 0 to 45.43; mean = 6.13 ± 10.39 SD) (Appendix 1 and Table 1), which motivated the analyses of each generation separately. Despite that, the coefficient of variation of abundance (both generations: from 0.75 to 4.36, mean =  $1.86 \pm 0.90$  SD) was largely similar between generations (0.52–2.65, mean =  $1.53 \pm 0.59$  SD for the spring generation, and 0.61-3.46, mean =  $1.62 \pm 0.73$  SD for the summer generation) (Appendix 1 and Table 1).

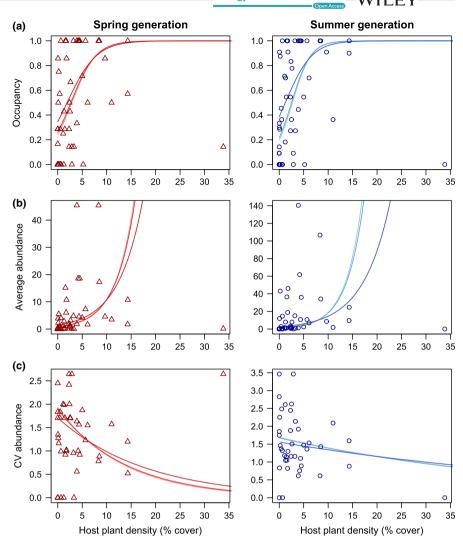
Results for the models explaining variation in occupancy among patches were generally similar between generations (Table 2 and Figure 3a). For both generations, the most important variables were patch area, connectivity, grazing frequency, and host plant density which all were positively related with occupancy and had relative importance = 1. Solar irradiance (mean and SD) also had a high relative importance in the models, especially in the summer generation (importance > 0.8), and revealed that patches with a higher and more variable microclimate were occupied more frequently. The interaction between connectivity and host plant density (importance = 0.83 and 1 for spring and summer generations, respectively) implies that, while occupancy generally increased with the density of host plants or with connectivity, host plant density had the largest effect in the most isolated patches, and conversely, connectivity had the largest effect on occupancy in patches with a low density of hosts (Figure 3a). Averaged parameter estimates showed that the effect of patch area (coefficient = 0. 61 and 0.51 for spring and summer generations), host plant density (0.85 and 0.72), and the interaction between host plant density and connectivity (-0.70 and -0.84) largely exceeded that of other variables (all other absolute coefficients < 0.5).

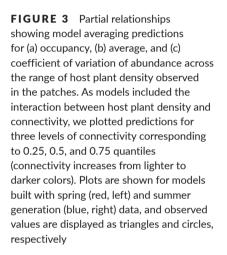
For average abundance, results from model averaging showed highly similar responses for both generations (Table 2 and Figure 3b). The most important variables explaining average abundance were patch area, connectivity, and host plant density (relative importance in all cases > 0.9). The interaction between connectivity and host plant density had also a high importance in the models (>0.8). As for occupancy, average abundance increased with patch area and host plant density or connectivity. Again, the interaction between the latter two revealed that highly connected patches were less sensitive to host density (Figure 3b). Grazing intensity had an intermediately high importance (variable importance = 0.58 and 0.32 for spring and summer generations, respectively), indicating that grazing allowed patches to sustain

**TABLE 2** Ninety-five percent model-averaged coefficients ( $\pm SE$ ) and variable importance from linear models explaining for each generation (a) occupancy, (b) average, and (c) coefficient of variation of abundance per patch. For grazing regime, the "never" category is taken as reference. Variable importance > 0.5 is highlighted in bold font, and the three highest absolute coefficient value for each model is displayed in italic font

	Spring genera	Spring generation			Summer generation		
Variables	Estimate	SE	Importance	Estimate	SE	Importance	
(a) Occupancy							
Area	0.621	0.181	1.000	0.512	0.113	1.000	
Connectivity	0.255	1.170	1.000	0.393	0.724	1.000	
Grazing (sometimes)	0.405	0.127	1.000	0.193	0.057	1.000	
Grazing (always)	0.244	0.141		0.136	0.062		
Host plant density	0.852	0.146	1.000	0.717	0.103	1.000	
Solar irradiance (mean)	0.049	0.074	0.453	0.121	0.040	1.000	
Solar irradiance (SD)	0.007	0.049	0.252	0.072	0.055	0.823	
Connectivity:Grazing (sometimes)	0.122	0.353	0.725	0.050	0.156	0.324	
Connectivity:Grazing (always)	0.400	0.515		0.120	0.250		
Connectivity:Host plant density	-0.697	0.431	0.828	-0.839	0.225	1.000	
Connectivity:Solar irradiance (mean)	-0.206	0.885	0.105	-0.239	0.626	0.266	
Connectivity:Solar irradiance (SD)	0.020	0.084	0.075	0.032	0.067	0.293	
(b) Average abundance							
Area	0.314	0.168	0.905	0.441	0.127	1.000	
Connectivity	0.121	1.140	1.000	0.200	0.936	1.000	
Grazing (sometimes)	0.252	0.290	0.578	0.105	0.202	0.315	
Grazing (always)	0.102	0.223		0.063	0.170		
Host plant density	0.901	0.264	1.000	0.871	0.242	1.000	
Solar irradiance (mean)	0.001	0.065	0.179	-0.007	0.063	0.184	
Solar irradiance (SD)	-0.017	0.086	0.206	0.012	0.076	0.211	
Connectivity:Grazing (sometimes)	0.208	0.433	0.227	0.125	0.322	0.159	
Connectivity:Grazing (always)	0.300	0.602		0.235	0.578		
Connectivity:Host plant density	-0.939	0.570	0.809	-1.052	0.512	0.866	
Connectivity:Solar irradiance (mean)	0.036	0.613	0.025	0.015	0.484	0.021	
Connectivity:Solar irradiance (SD)	0.006	0.064	0.033	0.005	0.056	0.033	
(c) CV abundance							
Area	-0.425	0.131	1.000	-0.268	0.202	0.779	
Connectivity	-0.036	1.458	1.000	-0.112	0.535	0.522	
Grazing (sometimes)	-0.335	0.278	0.724	-0.243	0.333	0.441	
Grazing (always)	-0.407	0.312		-0.240	0.331		
Host plant density	-1.126	0.263	1.000	-0.174	0.211	0.569	
Solar irradiance (mean)	-0.013	0.082	0.243	-0.004	0.081	0.188	
Solar irradiance (SD)	-0.011	0.067	0.168	-0.162	0.204	0.600	
Connectivity:Grazing (sometimes)	-0.014	0.142	0.024	-0.013	0.238	0.026	
Connectivity:Grazing (always)	-0.016	0.142	0.021	-0.031	0.330	0.020	
Connectivity:Host plant density	1.272	0.357	0.987	0.061	0.195	0.131	
Connectivity:Solar irradiance (mean)	-0.321	1.405	0.079	0.005	0.313	0.006	
Connectivity:Solar irradiance (SD)	0.001	0.032	0.012	-0.026	0.130	0.077	

a higher population size on average compared to nongrazed sites. All other variables or interactions had a relative importance between 0.03 and 0.23. Parameter estimates pointed to a greater effect of host plant density (coefficient = 0.90 and 0.87 for spring and summer generations), especially in interaction with connectivity (-0.94 and -1.05), compared to other variables (all other absolute coefficients < 0.5).





For the variability in abundance among years, models led to some notably different responses between generations (Table 2). In the spring generation, the variability in abundance tended to be reduced in large well-connected patches with a high host plant density (relative importance always = 1) and regularly grazed (0.72), although again the effect of plant density was reduced in highly connected patches (relative importance for interaction term = 0.99, Figure 3c). Model-averaged coefficients revealed that host plant density, alone (coefficient = -1.13) and in interaction with connectivity (coefficient = 1.27), had the highest effects on site variability during first generation. In contrast, the relative importance was more evenly distributed among variables in the second-generation model. Only patch area had a strong negative effect on abundance variability in the averaged model (importance = 0.78), followed by the standard deviation of solar irradiance (0.60) and host plant density (0.57) which both decreased the variability of abundance as they increased. Similarly, averaged estimates remained limited, with only area and grazing having a negative effect on site variability with a coefficient > 0.2. The full set of models used for model averaging is given in Appendix 2.

#### 4 | DISCUSSION

Although limiting factors may change in space and time (Lawson, Bennie, Thomas, Hodgson, & Wilson, 2012), understanding the determinants of distribution and population dynamics at species range margins is fundamental for our ability to predict biodiversity responses to climate change. We demonstrated that the occupancy, abundance, and population variability at the northern range margin of the butterfly P. armoricanus are mainly driven by patch area, connectivity, and host plant density, while solar irradiance only had an impact on patch occupancy. Large and well-connected habitat patches tended to be more often occupied, to have on average larger and more stable populations, and to display a higher abundance. The strong and consistent effects of patch area and connectivity show that the spatial configuration of habitat is the most important factor for population persistence at the climatic range margin of this butterfly. This pattern is congruent with metapopulation theory which predicts that occupancy and turnover rate are driven by patch area and site isolation (Hanski, 1994, 1998).

WILEY\_Ecology and Evolution

Several studies have highlighted the importance of microclimatic conditions for invertebrate populations, especially for populations near the climatic limits of the species (e.g., Bennie et al., 2013; Turlure, Choutt, Baguette, & Van Dyck, 2009; Wilson, Davies, & Thomas, 2010). We found that patches that received a high solar irradiancepresumably reflecting a warmer microclimate-were more frequently occupied than less sunny habitat patches, confirming previous findings that microclimate is an important aspect of habitat quality for this species (Eilers et al., 2013). In contrast, the effect of solar irradiance on abundance and population variability was negligible compared to other factors. Hence, our results suggest that some otherwise suitable habitat patches have a too cold microclimate to allow for persistence of P. armoricanus populations. On the other hand, where P. armoricanus is present, a higher solar irradiance does not result in larger populations. This could partly be because our surrogate for microclimate, that is, solar irradiance, only reflects the potential microclimate resulting from topography. The realized local temperature conditions can also be influenced by various factors such as vegetation cover or the surrounding landscape (Suggitt et al., 2011). A previous study found that the availability of host plants situated in a warm microclimate could predict local population sizes (Eilers et al., 2013). With warmer global temperatures, it is possible that this restriction to sites with a warm microclimate can be relaxed, and allow for colonization of a wider range of habitats at a regional scale, as has been observed for other butterflies at the climatic margins of their distributions (Pateman, Hill, Roy, Fox, & Thomas, 2012; Pateman, Thomas, Hayward, & Hill, 2016; Wilson et al., 2010). So far, however, this has not been observed for P. armoricanus in our study region.

The density of larval host plants, F. vulgaris and H. nummularium, appeared as a major driver of occupancy, abundance, and population variability. Larval host plants have long been known to play a vital role in the dynamics of butterfly populations (Hanski & Singer, 2001; Koh et al., 2004). As such, host plants' dynamics and distribution can strongly affect butterflies' response to climate change (Araújo & Luoto, 2007) and predicted range shifts by host plants could directly lead to changes or reductions in the distribution of herbivorous insects (Romo, Garcia-Barros, Marquez, Moreno, & Real, 2014; Romo, Silvestre, & Munguira, 2015). On the contrary, a mismatched response between a butterfly and its host may make it unable to track climate change (Pelini et al., 2009; Schweiger et al., 2008), even if host switching has also been documented (Pateman et al., 2012; Thomas et al., 2001). Our results suggest that host plant availability is one of the most important factors limiting the local abundance and landscape-scale distribution of P. armoricanus. However, at a larger spatial scale, the current distribution of P. armoricanus in Sweden is not limited by the distribution of its host plants, because both F. vulgaris and H. nummularium are known to occur much further north (Hultén, 1971). This leads us to believe that a potential expansion by the species to habitat further north is restricted by either microclimatic favorability or connectivity.

Connectivity had a particularly high importance in almost all models, especially in interaction with host plant density. Considering the effect of connectivity alone, our results reflect the fact that sites that are spatially isolated and surrounded by low-abundance sites have

generally a lower and more variable abundance and are less often occupied. This suggests that the current distribution of P. armoricanus could be constrained by a too high isolation of habitat patches further north, hence preventing their colonization due to limited dispersal. The effect of the interaction between host plant density and connectivity is also interesting; it reveals that the positive effect of host plants density on abundance or occupancy-and to a lesser extent site stabilitytends to decrease as connectivity increases. It can likely be interpreted as a rescue effect (Brown & Kodric-Brown, 1977) which allows lowquality sites to be sustained by migration from surrounding patches and reduced population fluctuations in well-connected patches. This property of metapopulation dynamics favors the persistence of a local population by decreasing extinction probability or by supporting population size through regular immigration events, even if the local conditions are suboptimal (Gonzalez, Lawton, Gilbert, Blackburn, & Evans-Freke, 1998; Gotelli, 1991). Because patches are separated by unsuitable intensively managed fields, the isolation of habitats is in the present case strongly driven by agricultural practices which determine the connectivity of the habitat network and will in this respect be key to the long-term persistence of this P. armoricanus population.

Our results challenge the still-common view that latitudinal edges are limited purely by climatic factors (Pearson & Dawson, 2003; Woodward, 1990), but concurs with modern niche theory that assumes actual distributional limits to be formed by an interaction between abiotic factors (fundamental niche, mainly climate), biotic interactions (here larval host), and dispersal (Soberón, 2007; Soberon & Nakamura, 2009). Better understanding of the actual position of this population relative to the species niche could be gained by investigating more closely its climatic tolerance, for example experimentally, or by comparing processes acting in various parts of its range (Lawson et al., 2012). In conclusion, it appears that the regional distribution and abundance of the northernmost population of P. armoricanus are mostly dependent on the availability of habitat patches with a high density of its larval host plants and on its capacity to disperse between such habitat patches. An effective conservation management strategy for this species should thus act both at the patch and landscape scales. First, habitat guality of already suitable patches must be maintained by ensuring the continuation of extensive grazing practices that provide an adequate vegetation structure and density of host plants. Second, the network of habitat patches must be kept dense enough to allow the long-term metapopulation viability. More generally, the persistence of many species in the face of climate change will be fundamentally linked to the maintenance of an efficient network of habitats (Hodgson, Thomas, Wintle, & Moilanen, 2009). It can be achieved by preserving the existent connectivity between habitats, but also by creating or restoring habitats to facilitate range shifts across a fragmented landscape (Hodgson, Wallis, Krishna, Cornell, & Isaac, 2016; Hodgson et al., 2011). In addition, enhancing local habitat quality at climatic range margins has been shown to be an effective alternative strategy to facilitate species expansion under climate change, as it secures vulnerable marginal populations and increases the pool of potential migrants (Lawson et al., 2012). Conservation planning should also take into account the current and future properties of the landscape matrix to assist species in tracking their favorable

**Ecology and Evolution** 

environmental conditions (Pearson & Dawson, 2005). Moreover, efficient actions should ideally consider the potential responses of all interacting species in an ecosystem (Walther, 2010), which makes management-assisted climate change mitigation challenging. However, maintaining the potential of species to respond effectively to the ongoing human-induced changes is essential for their persistence, especially for populations located at the margins of species distributions as, being generally located in suboptimal environmental conditions, they naturally have a higher extinction risk (Lesica & Allendorf, 1995).

## ACKNOWLEDGMENTS

Harriet Arnberg, Erik Cronvall, Åse Dannestam, Silke Eilers, Stefan Jarl, Per Karlsson, Anna Knöppel, Sara Leinerud, Mikael Molander, Josefin Olsson, Janielle Porter, Cecilia Ronnås, and Annika Söderman have assisted with the butterfly monitoring over the years. Thomas Ranius and two anonymous reviewers gave valuable comments on the manuscript. The study has been funded by the Swedish Research Council (contract 621-2010-5589), Signhild Engkvist Foundation, Lunds Djurskyddsfond, the Royal Swedish Academy of Agriculture and Forestry, and a strategic grant from the Department of Ecology SLU, to EÖ.

#### CONFLICT OF INTEREST

None declared.

#### REFERENCES

- Araújo, M. B., & Luoto, M. (2007). The importance of biotic interactions for modelling species distributions under climate change. *Global Ecology* and Biogeography, 16, 743–753.
- Barton, K. (2013) MuMIn: Multi-model inference. R package version 1.9.13. Retrieved from http://cran.r-project.org/package=MuMIn.
- Bennie, J., Hodgson, J. A., Lawson, C. R., Holloway, C. T. R., Roy, D. B., Brereton, T., ... Wilson, R. J. (2013). Range expansion through fragmented landscapes under a variable climate. *Ecology Letters*, 16, 921–929.
- Brown, J. H. (1984). On the relationship between abundance and distribution of species. *The American Naturalist*, 124, 255–279.
- Brown, J. H., & Kodric-Brown, A. (1977). Turnover rates in insular biogeography–Effect of immigration on extinction. *Ecology*, 58, 445–449.
- Brussard, P. F. (1984). Geographic patterns and environmental gradients: The central-marginal model in Drosophila revisited. Annual Review of Ecology and Systematics, 15, 25–64.
- Burnham, K. P., & Anderson, D. R. (2002). Model selection and multi-model inference: A practical information-theoretic approach. New York, NY: Springer-Verlag.
- Burrows, M. T., Schoeman, D. S., Richardson, A. J., Molinos, J. G., Hoffmann, A., Buckley, L. B., ... Poloczanska, E. S. (2014). Geographical limits to speciesrange shifts are suggested by climate velocity. *Nature*, 507, 492–495.
- Chen, I. C., Hill, J. K., Ohlemuller, R., Roy, D. B., & Thomas, C. D. (2011). Rapid range shifts of species associated with high levels of climate warming. *Science*, 333, 1024–1026.
- Christensen, P. R. (2000). The effects of grazing on the butterfly fauna in Denmark. PhD thesis, University of Aarhus, Aarhus, Denmark.
- Eilers, S., Pettersson, L. B., & Öckinger, E. (2013). Micro-climate determines oviposition site selection and abundance in the butterfly *Pyrgus armoricanus* at its northern range margin. *Ecological Entomology*, 38, 183–192.
- Eliasson, C., Ryrholm, N., Gärdenfors, U., Holmer, M., & Jilg, K. (2005). Dagfjärilar/Hesperiidae-Nymphalidae: Denna Volym Omfattar Samtliga

Nordiska Arter (Nationalnyckeln Till Sveriges Flora Och Fauna). ArtDatabanken, SLU, Uppsala, Sweden.

- Gillings, S., Balmer, D. E., & Fuller, R. J. (2015). Directionality of recent bird distribution shifts and climate change in Great Britain. *Global Change Biology*, 21, 2155–2168.
- Gilman, S. E., Urban, M. C., Tewksbury, J., Gilchrist, G. W., & Holt, R. D. (2010). A framework for community interactions under climate change. *Trends in Ecology & Evolution*, 25, 325–331.
- Gonzalez, A., Lawton, J. H., Gilbert, F. S., Blackburn, T. M., & Evans-Freke, I. I. (1998). Metapopulation dynamics, abundance, and distribution in a microecosystem. *Science*, 281, 2045–2047.
- Gotelli, N. J. (1991). Metapopulation models—The rescue effect, the propagule rain, and the core-satellite hypothesis. *The American Naturalist*, 138, 768–776.
- Hampe, A., & Petit, R. J. (2005). Conserving biodiversity under climate change: The rear edge matters. *Ecology Letters*, 8, 461–467.
- Hanski, I. (1994). A practical model of metapopulation dynamics. Journal of Animal Ecology, 63, 151–162.
- Hanski, I. (1998). Metapopulation dynamics. Nature, 396, 41-49.
- Hanski, I. (1999). Metapopulation ecology. Oxford: Oxford University Press.
- Hanski, I., & Singer, M. C. (2001). Extinction-colonization dynamics and host-plant choice in butterfly metapopulations. *The American Naturalist*, 158, 341–353.
- Hardie, D. C., & Hutchings, J. A. (2010). Evolutionary ecology at the extremes of species' ranges. *Environmental Reviews*, 18, 1–20.
- Hickling, R., Roy, D. B., Hill, J. K., Fox, R., & Thomas, C. D. (2006). The distributions of a wide range of taxonomic groups are expanding polewards. *Global Change Biology*, 12, 450–455.
- Hodgson, J. A., Thomas, C. D., Cinderby, S., Cambridge, H., Evans, P., & Hill, J. K. (2011). Habitat re-creation strategies for promoting adaptation of species to climate change. *Conservation Letters*, 4, 289–297.
- Hodgson, J. A., Thomas, C. D., Wintle, B. A., & Moilanen, A. (2009). Climate change, connectivity and conservation decision making: Back to basics. *Journal of Applied Ecology*, 46, 964–969.
- Hodgson, J. A., Wallis, D. W., Krishna, R., Cornell, S. J., & Isaac, N. (2016). How to manipulate landscapes to improve the potential for range expansion. *Methods in Ecology and Evolution*, in press, doi: 10.1111/2041-210x.12614
- Hultén, E. (1971). Atlas Över Växternas Utbredning I Norden. Stockholm: Generalstabens litografiska anstalts förlag.
- Jetz, W., Wilcove, D. S., & Dobson, A. P. (2007). Projected impacts of climate and land-use change on the global diversity of birds. *PLoS Biology*, 5, e157.
- Kleijn, D., Schekkerman, H., Dimmers, W. J., Van Kats, R. J. M., Melman, D., & Teunissen, W. A. (2010). Adverse effects of agricultural intensification and climate change on breeding habitat quality of Black-tailed Godwits Limosa I. limosa in the Netherlands. *Ibis*, 152, 475–486.
- Koh, L. P., Dunn, R. R., Sodhi, N. S., Colwell, R. K., Proctor, H. C., & Smith, V. S. (2004). Species coextinctions and the biodiversity crisis. *Science*, 305, 1632–1634.
- Kruess, A., & Tscharntke, T. (2002). Grazing intensity and the diversity of grasshoppers, butterflies, and trap-nesting bees and wasps. *Conservation Biology*, 16, 1570–1580.
- Kudrna, O., Harpke, A., Lux, K., Pennerstorfer, J., Schweiger, O., Settele, J., & Wiemers, M. (2011). Distribution atlas of butterflies in Europe. Halle, Germany: GfS.
- Lantmäteriet (2015). Product description: GSD-Elevation data, Grid 2+. Document version 2.2. Retrieved from https://www.lantmateriet.se/sv/ Kartor-och-geografisk-information/Hojddata/GSD-Hojddata-grid-2/#.
- Lawson, C. R., Bennie, J., Hodgson, J. A., Thomas, C. D., & Wilson, R. J. (2014). Topographic microclimates drive microhabitat associations at the range margin of a butterfly. *Ecography*, 37, 732–740.
- Lawson, C. R., Bennie, J. J., Thomas, C. D., Hodgson, J. A., & Wilson, R. J. (2012). Local and landscape management of an expanding range margin under climate change. *Journal of Applied Ecology*, 49, 552–561.

WILEY\_Ecology and Evolution

- Lesica, P., & Allendorf, F. W. (1995). When are peripheral populations valuable for conservation? *Conservation Biology*, *9*, 753–760.
- Louthan, A. M., Doak, D. F., & Angert, A. L. (2015). Where and when do species interactions set range limits? *Trends in Ecology & Evolution*, 30, 780–792.
- MacKenzie, D. I., Nichols, J. D., Lachman, G. B., Droege, S., Andrew Royle, J., & Langtimm, C. A. (2002). Estimating site occupancy rates when detection probabilities are less than one. *Ecology*, 83, 2248–2255.
- Nicolè, F., Dahlgren, J. P., Vivat, A., Till-Bottraud, I., & Ehrlén, J. (2011). Interdependent effects of habitat quality and climate on population growth of an endangered plant. *Journal of Ecology*, 99, 1211–1218.
- Nordström, F., Opheim, M., & Valle, K. (1955). De Fennoskandiska dagfjärilarnas utbredning; Lepidoptera diurna (Rhopalocera & Hesperioidea). Lund, Sweden: C.W.K. Gleerup.
- Öckinger, E. (2006). Possible metapopulation structure of the threatened butterfly Pyrgus armoricanus in Sweden. Journal of Insect Conservation, 10, 43–51.
- Öckinger, E. (2007). Backvisslaren Pyrgus armoricanus (Lepidoptera: Hesperiidae)—en förbisedd klimatvinnare? Entomologisk Tidskrift, 128, 9-17.
- Ojanen, S. P., Nieminen, M., Meyke, E., Poyry, J., & Hanski, I. (2013). Longterm metapopulation study of the Glanville fritillary butterfly (*Melitaea cinxia*): Survey methods, data management, and long-term population trends. *Ecology and Evolution*, *3*, 3713–3737.
- Opdam, P., & Wascher, D. (2004). Climate change meets habitat fragmentation: Linking landscape and biogeographical scale levels in research and conservation. *Biological Conservation*, 117, 285–297.
- Parmesan, C., Ryrholm, N., Stefanescu, C., Hill, J. K., Thomas, C. D., Descimon, H., ... Warren, M. (1999). Poleward shifts in geographical ranges of butterfly species associated with regional warming. *Nature*, 399, 579–583.
- Pateman, R. M., Hill, J. K., Roy, D. B., Fox, R., & Thomas, C. D. (2012). Temperature-dependent alterations in host use drive rapid range expansion in a butterfly. *Science*, 336, 1028–1030.
- Pateman, R. M., Thomas, C. D., Hayward, S. A., & Hill, J. K. (2016). Macroand microclimatic interactions can drive variation in species' habitat associations. *Global Change Biology*, 22, 556–566.
- Pearson, R. G., & Dawson, T. P. (2003). Predicting the impacts of climate change on the distribution of species: Are bioclimate envelope models useful? *Global Ecology and Biogeography*, 12, 361–371.
- Pearson, R. G., & Dawson, T. P. (2005). Long-distance plant dispersal and habitat fragmentation: Identifying conservation targets for spatial landscape planning under climate change. *Biological Conservation*, 123, 389–401.
- Pelini, S. L., Dzurisin, J. D., Prior, K. M., Williams, C. M., Marsico, T. D., Sinclair, B. J., & Hellmann, J. J. (2009). Translocation experiments with butterflies reveal limits to enhancement of poleward populations under climate change. Proceedings of the National Academy of Sciences of the United States of America, 106, 11160–11165.
- R Development Core Team (2015) *R*: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. URL: http://www.R-project.org/.
- Romo, H., Garcia-Barros, E., Marquez, A. L., Moreno, J. C., & Real, R. (2014). Effects of climate change on the distribution of ecologically interacting species: Butterflies and their main food plants in Spain. *Ecography*, 37, 1063–1072.
- Romo, H., Silvestre, M., & Munguira, M. L. (2015). Potential distribution models and the effect of climatic change on the distribution of *Phengaris nausithous* considering its food plant and host ants. *Journal of Insect Conservation*, 19, 1101–1118.

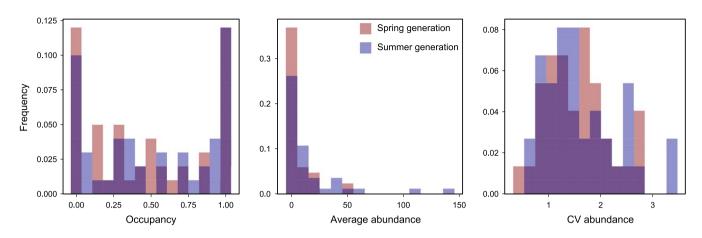
- Roy, D. B., & Thomas, J. A. (2003). Seasonal variation in the niche, habitat availability and population fluctuations of a bivoltine thermophilous insect near its range margin. *Oecologia*, 134, 439–444.
- Sagarin, R. D., & Gaines, S. D. (2002). The 'abundant centre' distribution: To what extent is it a biogeographical rule? *Ecology Letters*, 5, 137–147.
- Schweiger, O., Settele, J., Kudrna, O., Klotz, S., & Kuhn, I. (2008). Climate change can cause spatial mismatch of trophically interacting species. *Ecology*, 89, 3472–3479.
- Seabrook, L., McAlpine, C., Rhodes, J., Baxter, G., Bradley, A., & Lunney, D. (2014). Determining range edges: Habitat quality, climate or climate extremes? *Diversity and Distributions*, 20, 95–106.
- Sexton, J. P., McIntyre, P. J., Angert, A. L., & Rice, K. J. (2009). Evolution and ecology of species range limits. *Annual Review of Ecology, Evolution, and Systematics*, 40, 415–436.
- Smale, D. A., & Wernberg, T. (2013). Extreme climatic event drives range contraction of a habitat-forming species. Proceedings of the Royal Society B: Biological Sciences, 280, 20122829.
- Soberón, J. (2007). Grinnellian and Eltonian niches and geographic distributions of species. *Ecology Letters*, 10, 1115–1123.
- Soberon, J., & Nakamura, M. (2009). Niches and distributional areas: Concepts, methods, and assumptions. Proceedings of the National Academy of Sciences of the United States of America, 106, 19644–19650.
- Suggitt, A. J., Gillingham, P. K., Hill, J. K., Huntley, B., Kunin, W. E., Roy, D. B., & Thomas, C. D. (2011). Habitat microclimates drive fine-scale variation in extreme temperatures. *Oikos*, 120, 1–8.
- Thomas, J. A. (1993). Holocene climate changes and warm man-made refugia may explain why a sixth of British butterflies possess unnatural early-successional habitats. *Ecography*, 16, 278–284.
- Thomas, C. D., Bodsworth, E. J., Wilson, R. J., Simmons, A. D., Davies, Z. G., Musche, M., & Conradt, L. (2001). Ecological and evolutionary processes at expanding range margins. *Nature*, 411, 577–581.
- Thuiller, W., Albert, C., Araujo, M. B., Berry, P. M., Cabeza, M., Guisan, A., ... Zimmermann, N. E. (2008). Predicting global change impacts on plant species' distributions: Future challenges. *Perspectives in Plant Ecology Evolution and Systematics*, 9, 137–152.
- Turlure, C., Choutt, J., Baguette, M., & Van Dyck, H. (2009). Microclimatic buffering and resource-based habitat in a glacial relict butterfly: Significance for conservation under climate change. *Global Change Biology*, *16*, 1883–1893.
- Van der Putten, W. H., Macel, M., & Visser, M. E. (2010). Predicting species distribution and abundance responses to climate change: Why it is essential to include biotic interactions across trophic levels. *Philosophical Transactions of the Royal Society B: Biological Sciences*, 365, 2025–2034.
- Walther, G. R. (2010). Community and ecosystem responses to recent climate change. *Philosophical Transactions of the Royal Society B: Biological Sciences*, 365, 2019–2024.
- Wilson, R. J., Davies, Z. G., & Thomas, C. D. (2010). Linking habitat use to range expansion rates in fragmented landscapes: A metapopulation approach. *Ecography*, 33, 73–82.
- Woodward, F. I. (1990). The impact of low-temperatures in controlling the geographical-distribution of plants. *Philosophical Transactions of the Royal Society B: Biological Sciences*, 326, 585–593.

How to cite this article: Fourcade, Y. and Öckinger, E. (2017), Host plant density and patch isolation drive occupancy and abundance at a butterfly's northern range margin. Ecology and Evolution, 7: 331–345. doi: 10.1002/ece3.2597

VILEY

# **APPENDIX 1**

Frequency histograms of occupancy, average abundance, and coefficient of variation of abundance, for the spring (red) and summer (blue) generations. Overlapping areas appear in purple.



#### **APPENDIX 2**

Models ranked by  $AIC_{c}$ , explaining (a) occupancy, (b) average, and (c) coefficient of variation of abundance, for each generation. Only models used in model averaging are shown, that is, those whose cumulative weight was < 95%. Full models (including all variables and interactions) and null models (including only intercept) are also shown for comparison. Variables are abbreviated as follows: A: area; C: connectivity; HD: host plant density; G: grazing intensity; SRM: mean solar irradiance; SRSD: standard deviation of solar irradiance.

/ariables	df	logLik	AIC <sub>c</sub>	$\Delta AIC_{c}$	ω
a) occupancy					
Spring generation					
A + C + G + HD + C:G + C:HD	9	-79.714	181.929	0.000	.237
A + C + G + HD + SRM + C:G + C:HD	10	-78.523	182.687	0.758	.163
A + C + G + HD + C:G	8	-82.347	184.206	2.277	.076
A + C + G + HD + C:HD	7	-83.944	184.554	2.625	.064
A + C + G + HD + SRM + C:G	9	-81.238	184.976	3.047	.052
A + C + G + HD + SRSD + C:G + C:HD	10	-79.693	185.028	3.099	.050
A + C + G + HD + SRSD + C:HD + C:SRSD	9	-81.427	185.354	3.425	.043
A + C + G + HD + SRM + C:G + C:HD + C:SRM	11	-78.269	185.486	3.557	.040
A + C + G + HD + SRM + SRSD + C:G + C:HD	11	-78.485	185.916	3.988	.032
A + C + G + HD + SRM + C:HD	8	-83.303	186.117	4.188	.029
A + C + G + HD + SRSD + C:HD	8	-83.304	186.120	4.191	.029
A + C + G + HD + SRM + C:HD + C:SRM	9	-81.829	186.159	4.230	.029
A + C + G + HD + SRM + SRSD + C:HD + C:SRSD	10	-80.279	186.199	4.270	.028
A + C + G + HD + SRM + SRSD + C:HD	9	-81.948	186.395	4.466	.025
A + C + G + HD + SRM + C:G + C:SRM	10	-80.760	187.161	5.232	.017
A + C + G + HD + SRSD + C:G	9	-82.347	187.193	5.264	.017
Full model	13	-78.074	192.259	10.330	.001
Null model	1	-157.873	317.829	135.900	.000
Summer generation					
A + C + G + HD + SRM + SRSD + C:HD	9	-110.154	242.809	0.000	.251

342

WILEY\_Ecology and Evolution

Variables df logLik   A + C + G + HD + SRM + SRSD + C:HD + C:SRSD 10 -108.751   A + C + G + HD + SRM + SRSD + C:HD + C:SRM 10 -109.388   A + C + G + HD + SRM + SRSD + C:HD + C:SRM 10 -109.388	AIC <sub>c</sub> 243.144 244.416 244.475	ΔΑΙC <sub>c</sub> 0.335	ω .212
A + C + G + HD + SRM + SRSD + C:HD + C:SRM 10 -109.388	244.416		.212
		1 (00	.212
	244.475	1.608	.112
A + C + G + HD + SRM + SRSD + C:G + C:HD 11 -107.764		1.667	.109
A + C + G + HD + SRM + C:G + C:HD 10 -109.501	244.644	1.835	.100
A + C + G + HD + SRM + C:G + C:HD + C:SRM 11 -108.701	246.349	3.540	.043
A + C + G + HD + SRM + SRSD + C:HD + C:SRM + C:SRSD 11 -108.722	246.392	3.584	.042
A + C + G + HD + SRM + SRSD + C:G + C:HD + C:SRM 12 -107.277	246.987	4.179	.031
A + C + G + HD + SRM + C:HD + C:SRM 9 -112.458	247.415	4.606	.025
A + C + G + HD + SRM + SRSD + C:G + C:HD + C:SRSD 12 -107.519	247.471	4.662	.024
Full model 13 -107.274   Null model 13 -204.574	250.660	7.851	.005
Null model 1 -234.571	471.225	228.416	.000
(b) Average abundance Spring generation			
$A + C + HD + C:HD \qquad 6 \qquad -67.907$	150.284	0.000	.234
A + C + G + HD + C:HD 8 -65.099	150.697	0.413	.190
A + C + G + HD + C:HD A + C + G + HD + C:G 9 -64.114	152.035	1.751	.098
A + C + HD + SRSD + C:HD 7 -67.784	152.962	2.678	.061
A + C + HD + SRM + C:HD 7 -67.889	153.171	2.887	.055
A + C + G + HD + SRSD + C:HD 9 -65.085	153.976	3.692	.037
A + C + G + HD + SRM + C:HD 9 -65.088	153.983	3.699	.037
C + G + HD + C:HD 7 -68.541	154.475	4.191	.029
A + C + G + HD + SRSD + C:G 10 -63.791	154.916	4.632	.023
A + C + G + HD + C:G + C:HD 10 -63.873	155.080	4.796	.021
C + G + HD + C:G 8 -67.333	155.166	4.882	.020
A + C + G + HD + SRM + C:G 10 -64.088	155.509	5.225	.017
A + C + HD + SRSD + C:HD + C:SRSD 8 -67.638	155.775	5.491	.015
A + C + HD + SRM + SRSD + C:HD 8 -67.784	156.067	5.783	.013
A + C + HD + SRM + C:HD + C:SRM 8 -67.791	156.081	5.797	.013
A + C + G + HD + SRSD + C:HD + C:SRSD 10 -64.632	156.598	6.314	.010
C + HD + C:HD 5 -72.652	157.019	6.735	.008
C + G + HD + SRSD + C:G 9 -66.654	157.115	6.831	.008
C + G + HD + SRSD + C:HD 8 -68.458   A + C + G + HD + SRM + SRSD + C:HD 10 -65.062	157.417	7.133	.007
A + C + G + HD + SRM + SRSD + C:HD 10 -65.062   C + G + HD + SRM + C:HD 8 -68.488	157.457 157.476	7.173 7.192	.008
A + C + G + HD + SRM + C:HD + C:SRM 10 -65.082	157.498	7.214	.006
C + G + HD + C:G + C:HD 9 -67.136	158.079	7.795	.005
A + C + G + HD + SRSD + C:G + C:HD 11 -63.494	158.092	7.808	.005
A + C + G + HD + SRM + C:G + C:SRM 11 -63.556	158.216	7.932	.004
C + G + HD + SRM + C:G 9 -67.328	158.462	8.178	.004
A + C + G + HD + SRSD + C:G + C:SRSD 11 -63.770	158.644	8.360	.004
A + C + G + HD + SRM + SRSD + C:G 11 -63.791	158.685	8.401	.004
A + C + G + HD + SRM + C:G + C:HD 11 -63.843	158.789	8.505	.003
C + G + HD + SRSD + C:HD + C:SRSD 9 -67.547	158.901	8.617	.003
C + HD + SRSD + C:HD 6 -72.284	159.038	8.754	.003
Null model 2 -79.305	162.927	12.643	.000

# **APPENDIX 2** (Continued)

/ariables	df	logLik	AIC <sub>c</sub>	$\Delta AIC_{c}$	ω
Full model	14	-62.529	169.211	18.927	.000
Summer generation					
A + C + HD + C:HD	6	-70.814	156.099	0.000	.380
A + C + HD + SRSD + C:HD	7	-70.633	158.660	2.561	.106
A + C + G + HD + C:HD	8	-69.166	158.831	2.732	.097
A + C + G + HD + C:G	9	-67.543	158.892	2.793	.094
A + C + HD + SRM + C:HD	7	-70.760	158.915	2.816	.093
A + C + HD + SRSD + C:HD + C:SRSD	8	-70.486	161.472	5.373	.026
A + C + G + HD + C:G + C:HD	10	-67.146	161.626	5.527	.024
A + C + G + HD + SRSD + C:HD	9	-68.920	161.645	5.546	.024
A + C + HD + SRM + SRSD + C:HD	8	-70.625	161.750	5.651	.023
A + C + G + HD + SRM + C:HD	9	-69.012	161.830	5.731	.022
A + C + HD + SRM + C:HD + C:SRM	8	-70.730	161.960	5.860	.020
A + C + G + HD + SRSD + C:G	10	-67.541	162.416	6.317	.016
A + C + G + HD + SRM + C:G	10	-67.543	162.419	6.320	.016
A + C + G + HD + SRSD + C:HD + C:SRSD	10	-68.621	164.575	8.476	.005
Null model	2	-85.588	175.492	19.393	.000
Full model	14	-66.340	176.834	20.735	.000
c) coefficient of variation of abundance					
Spring generation					
A + C + G + HD + C:HD	8	-1.678	24.689	0.000	.421
A + C + HD + C:HD	6	-5.920	26.737	2.048	.151
A + C + G + HD + SRM + C:HD	9	-1.513	27.949	3.260	.083
A + C + G + HD + SRSD + C:HD	9	-1.603	28.130	3.441	.075
A + C + G + HD + SRM + C:HD + C:SRM	10	-0.148	29.096	4.407	.047
A + C + HD + SRM + C:HD	7	-5.548	29.097	4.407	.046
A + C + HD + SRSD + C:HD	7	-5.838	29.676	4.986	.035
A + C + HD + SRM + C:HD + C:SRM	8	-4.865	31.063	6.374	.017
A + C + G + HD + SRM + SRSD + C:HD	10	-1.317	31.433	6.744	.014
A + C + G + HD + C:G	9	-3.409	31.740	7.051	.012
A + C + G + HD + SRSD + C:HD + C:SRSD	10	-1.526	31.851	7.162	.012
A + C + HD + SRM + SRSD + C:HD	8	-5.261	31.856	7.167	.012
A + C + G + HD + SRM + SRSD + C:HD + C:SRM	11	0.467	32.066	7.377	.011
A + C + G + HD + C:G + C:HD	10	-1.678	32.155	7.466	.010
Null model	2	-19.095	42.554	17.865	.000
Full model	14	1.620	44.760	20.071	.000
Summer generation					
A + SRSD	4	-17.826	44.864	0.000	.064
A + G + HD + SRSD	7	-13.711	45.155	0.292	.055
A + C + SRSD	5	-16.669	45.213	0.349	.054
A + HD + SRSD	5	-16.922	45.719	0.856	.042
A + G + HD	6	-15.639	45.988	1.124	.036
A + G + SRSD	6	-15.702	46.113	1.249	.034
A + C	4	-18.574	46.361	1.497	.030

## **APPENDIX 2** (Continued)

ables	df	logLik	AIC <sub>c</sub>	∆AIC <sub>c</sub>	ω
A + C + HD + C:HD	6	-15.853	46.415	1.551	.02
A + C + HD + SRSD	6	-16.036	46.781	1.917	.02
A + C + HD	5	-17.540	46.954	2.090	.02
A + HD	4	-18.970	47.152	2.288	.02
A + C + G + HD + C:HD	8	-13.104	47.174	2.310	.02
G + HD	5	-17.694	47.262	2.398	.0:
A + C + SRSD + C:SRSD	6	-16.310	47.330	2.466	.0:
A + SRM + SRSD	5	-17.808	47.491	2.627	.0
G + HD + SRSD	6	-16.411	47.532	2.668	.0
A + C + SRM + SRSD	6	-16.462	47.633	2.770	.0
A + C + G + SRSD	7	-14.959	47.651	2.787	.0
A + C + G + HD + SRSD	8	-13.386	47.738	2.875	.0
A + C + G + HD	7	-15.019	47.772	2.908	.0:
A	3	-20.587	47.879	3.015	.0:
G + SRSD	5	-18.026	47.927	3.064	.0
A + C + HD + SRSD + C:HD	7	-15.147	48.028	3.164	.0
A + C + HD + SRSD + C:SRSD	7	-15.191	48.115	3.251	.0
C + G + HD + C:HD	7	-15.236	48.206	3.342	.0
SRSD	3	-20.779	48.263	3.400	.0
A + G + HD + SRM + SRSD	8	-13.666	48.298	3.435	.0
A + HD + SRM + SRSD	6	-16.838	48.386	3.522	.0
A + C + G + HD + C:G	9	-12.141	48.711	3.847	.0
A + G + HD + SRM	7	-15.533	48.799	3.935	.0
С	3	-21.092	48.889	4.025	.0
A + C + G + HD + SRSD + C:HD	9	-12.250	48.928	4.064	.0
A + C + SRM	5	-18.567	49.009	4.145	.0
C + G + HD + C:G	8	-14.026	49.018	4.154	.0
A + C + G + HD + SRSD + C:SRSD	9	-12.313	49.054	4.190	.0
A + C + G	6	-17.209	49.127	4.263	.0
A + G + SRM + SRSD	7	-15.701	49.136	4.273	.0
A + G	5	-18.644	49.162	4.298	.0
A + C + HD + SRM + SRSD	7	-15.724	47.102	4.278	.0
	2				
Null model		-22.441	49.224	4.360	.0. 0.
	4	-20.041 -15.792	49.294	4.431	
A + C + HD + SRM + C:HD	7		49.316	4.453	0.
	3	-21.316	49.338	4.474	0.
C + G + HD G	6	-17.345	49.401 49.519	4.537 4.656	.0
	4	-20.154			.0
C + HD + C:HD	5	-18.832	49.539	4.675	.0
HD + SRSD	4	-20.170	49.553	4.689	.0
A + HD + SRM	5	-18.890	49.654	4.791	.00
A + SRM	4	-20.267	49.745	4.881	.00
A + C + HD + SRM	6	-17.531	49.771	4.907	.00

iables	df	logLik	AIC <sub>c</sub>	ΔAIC <sub>c</sub>	ω
C + G + SRSD	6	-17.579	49.869	5.005	.00
A + C + SRM + SRSD + C:SRSD	7	-16.126	49.986	5.122	.00
C + HD	4	-20.402	50.017	5.153	.00
A + C + HD + SRSD + C:HD + C:SRSD	8	-14.574	50.114	5.250	.00
C + G	5	-19.177	50.228	5.364	.00
A + C + G + SRSD + C:SRSD	8	-14.638	50.242	5.378	.00
C + G + HD + SRSD	7	-16.258	50.250	5.386	.00
C + G + HD + SRSD + C:SRSD	8	-14.706	50.378	5.514	.00
C + G + HD + SRSD + C:HD	8	-14.788	50.542	5.678	.00
G + HD + SRM + SRSD	7	-16.404	50.542	5.678	.00
C + SRSD + C:SRSD	5	-19.351	50.577	5.713	.0
A + C + G + HD + SRM + C:HD	9	-13.100	50.629	5.765	.0
A + C + HD + SRM + SRSD + C:HD	8	-14.836	50.638	5.774	.0
A + C + SRM + SRSD + C:SRM	7	-16.462	50.657	5.793	.0
A + C + G + SRM + SRSD	8	-14.879	50.723	5.859	.0
A + C + HD + SRM + SRSD + C:SRSD	8	-14.883	50.731	5.867	.0
G + SRM + SRSD	6	-18.018	50.745	5.881	.0
SRM + SRSD	4	-20.778	50.768	5.904	.0
A + C + G + HD + SRM + SRSD	9	-13.236	50.901	6.037	.0
A + C + G + HD + SRM	8	-15.013	50.991	6.128	.0
A + G + SRM	6	-18.176	51.062	6.198	.0
SRM	3	-22.187	51.079	6.215	.0
A + C + G + HD + SRSD + C:HD + C:SRSD	10	-11.472	51.091	6.228	.0
C + HD + SRSD	5	-19.621	51.118	6.254	.0
G + SRM	5	-19.709	51.292	6.428	.0
C + SRM	4	-21.079	51.371	6.507	.0
C + G + HD + SRM + C:HD	8	-15.236	51.438	6.574	.0
C + HD + SRSD + C:SRSD	6	-18.372	51.454	6.590	.0
A + C + SRM + C:SRM	6	-18.401	51.512	6.648	.0
C + G + SRSD + C:SRSD	7	-16.950	51.633	6.769	.0
HD + SRM	4	-21.246	51.704	6.840	.0
A + C + G + HD + C:G + C:HD	10	-11.799	51.746	6.882	.0
C + G + HD + SRSD + C:HD + C:SRSD	9	-13.666	51.760	6.896	.0
C + SRM + SRSD	5	-19.966	51.807	6.944	.0
A + C + G + HD + SRSD + C:G	10	-11.839	51.825	6.962	.0
C + G + HD + C:G + C:HD	9	-13.700	51.828	6.964	.0
C + HD + SRSD + C:HD	6	-18.591	51.891	7.027	.00
A + C + G + HD + SRM + C:G	10	-11.891	51.931	7.067	.00
A + C + G + SRM	7	-17.132	51.998	7.134	.00
Full model	14	-10.298	66.857	21.993	.00