

Habitat amount modulates the effect of patch isolation on host-parasitoid interactions

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Habitat amount and patch isolation are important determinants of biodiversity and ecosystem functioning. We studied the separate and interactive effects of these two components of habitat fragmentation on host-parasitoid interactions in a replicated landscape-scale study. We used trap-nesting solitary bees, wasps and their natural enemies as study system. We exposed trap nests in 30 tree patches in agricultural landscapes in northern Switzerland. Study sites were either (i) adjacent to forest (adjacent), (ii) distant from forest but connected through woody elements (connected) or (iii) distant from forest with no connecting woody elements (isolated). Independent of the three levels of isolation, the amount of woody habitat in the landscapes covered a gradient from 4 to 74%. Host and parasitoid species richness increased with the amount of habitat in the landscape and was strongly reduced at isolated compared to adjacent and connected sites. Loss of host species richness was 21% at isolated compared to non-isolated sites, whereas parasitoid species richness decreased by 68%, indicating that the higher trophic level was more adversely affected by isolation. Most importantly, habitat amount and isolation had a pronounced interactive effect on parasitism: while isolation resulted in a strong decrease in parasitism in landscapes with low habitat amount, this effect was mitigated by high habitat amount. These interactive effects were consistent across the three years of the study. The observed interplay between habitat amount and patch isolation may explain the often conflicting results in the habitat fragmentation literature and should be considered in future research on multitrophic communities and ecosystem functioning in fragmented landscapes.

Keywords: connectivity, habitat fragmentation, Hymenoptera, landscape ecology, multitrophic interactions, solitary bee

INTRODUCTION

Habitat fragmentation can affect the diversity of species and disrupt their interactions (Aizen et al., 2012; Hagen et al., 2012; Ferreira et al., 2013; Rösch et al., 2013), with consequences for ecosystem service provisioning (Kruess and Tscharntke, 1994; Staddon et al., 2010; Tylianakis, 2013; Schüepp et al., 2014a,b). However, our understanding of the impact of fragmentation on ecological communities and the functional consequences for ecosystems suffers from major gaps (Tscharntke et al., 2012). Fragmentation consists of habitat loss and fragmentation per se, the latter defined as the breakdown of core habitat into isolated patches (Fahrig, 2003). Separating the consequences of patch isolation from that of habitat loss in the landscape is a challenge in empirical studies (Tscharntke et al., 2012). Yet, such an approach may be key for advancing our understanding of biodiversity and ecosystem functioning in a fragmented world (Farwig et al., 2009; Hadley and Betts, 2011; Mortelliti et al., 2011; With and Pavuk, 2011; Herrera and Doblas-Miranda, 2013). In particular, potential interactive effects among habitat loss and isolation on trophic interactions remain largely unexplored (Herrera

and Doblas-Miranda, 2013). Visser et al. (2009) modeled such interactive effects for a simple host-parasitoid system, predicting interactions on parasitoid persistence, whereas parasitism should be primarily negatively affected by isolation, regardless of habitat amount. In contrast to these predictions, Fahrig (2013) posits that ecological responses at the patch level should be essentially driven by the amount of habitat within the landscape, irrespective of patch isolation and size (habitat amount hypothesis). Because species at higher positions in the food chain may be particularly vulnerable to habitat loss (Holt et al., 1999; Kruess and Tscharntke, 2000; Cagnolo et al., 2009; Valladares et al., 2012), species richness of parasitoids should increase strongly with the amount of habitat in a landscape, even if the size of the sampled habitat patch is constant (Fahrig, 2013). If higher parasitoid diversity leads to higher parasitism rate (e.g., Tylianakis et al., 2006; Fenoglio et al., 2012), negative effects of isolation on parasitism rate, as predicted by Visser et al. (2009), may be mitigated by higher habitat amount within the landscape. As a result, decline in parasitism rate with isolation should be more pronounced in simple, habitat-poor landscapes compared to more complex

landscapes with high habitat amount. Here, we investigated the separate and interactive effects of habitat amount and isolation on the trophic interactions of trap-nesting bees and wasps with their parasitoids over three years. In particular, we tested the following hypotheses:

- 1) Species richness and abundance in host-parasitoid communities increase with increasing habitat amount within the landscape and are lower in isolated compared to adjacent and connected patches. Patterns are more pronounced for the higher compared to the lower trophic level.
- 2) Parasitism is reduced in isolated patches and in habitat patches surrounded by low amount of habitat as a result of reduced parasitoid species richness and abundance.
- 3) High habitat amount mitigates adverse effects of patch isolation on host and parasitoid diversity and on parasitism.

MATERIALS AND METHODS

STUDY SITES AND EXPERIMENTAL DESIGN

The study was conducted on the Swiss Plateau between the cities of Bern, Solothurn and Fribourg, a region that is dominated by agriculture interspersed with forest patches. Thirty experimental sites were chosen over an area of 23×32 km varying in altitude between 465 and 705 m above sea level (**Figure 1**). The sites were selected according to the amount of woody habitat (forests, hedgerows and orchards) within a 500 m radius (ranging from 4 to 74% woody habitat). Circles of 500 m radius were chosen because (i) this scale is considered to roughly match the activity range of the studied organisms (e.g., Gathmann and Tscharntke, 2002; Zurbuchen et al., 2010) and (ii) they correspond to the rather fine "grain" of the landscape of that region, i.e., the small-scaled mosaic of woody and agricultural landscape elements. The second selection criterion was isolation of the sites from woody habitat (**Figure 2**):

- Ten sites were located adjacent to forest (adjacent);
- ten sites were located at a distance of 100–200 m from the next forest, but next to woody elements such as hedgerows or rows of single trees that partly filled the interspace between study site and forest (connected);
- ten sites were located at least 100 m away from any woody habitat (isolated).

We distinguished among adjacent and connected sites because of the structural properties of the adjoining woody habitat. Forest had a closed canopy and no gaps between the shaded areas, while the connecting small-sized woody elements had gaps of open habitat between trees and shrubs, as did our study sites. Information on woody habitats was derived from official digital land-use maps (vector25, swisstopo, Wabern) and verified using aerial photographs and field inspection. There was no statistical dependency between the percentage of woody habitat in a landscape sector and the level of isolation of a study site [$F_{(2, 27)} =$ 0.004, P = 0.99]. The distance to the nearest forest was similar among isolated and connected sites (isolated: 128.5 ± 31.7 m, connected: 147.7 ± 41.6 m). The 500 m-radius landscape sectors



FIGURE 1 | Localisation of the 30 study sites in the Swiss lowlands. Green: forest; gray: settlements; blue: open water. Map background Vector 200 ©Swisstopo 2014.

surrounding the study sites were located at least 750 m apart to minimize potential spatial autocorrelation. In order to standardize the habitat type in which the investigations were to be conducted, at each site we planted an 18 m-long row of young cherry trees on permanent grassland in spring 2008. Since then, the study sites were managed in a standardized manner (Schüepp et al., 2011).

In other words, within the 500 m landscape circles, we measured at adjacent, connected and isolated locations but on artificially introduced, standardized cherry tree rows as habitat patches.

STUDY SYSTEM

In mid-March 2008, four trap nests for solitary bees and wasps were set up in each habitat patch on wooden posts 1 m above ground. Two traps were placed at 6 m distance from each end of each experimentally established tree row. Trap-nests consisted of plastic cylinders filled with an average of 180 tubes of common reed (*Phragmites australis* L.), with diameters ranging from 2 to 10 mm and the same proportion of each diameter in every trap. Each year from 2008 to 2010 two of the four trap nests were removed in October to analyse wasp and bee communities, and two were left in the field to allow for local population



development over these three consecutive years. Removed trap nests were stored at 5°C from mid-October to mid-January, and occupied reed internodes were individually transferred into glass tubes. The glass tubes were left at room temperature (22°C) for the emergence of adult bees, wasps and their natural enemies (enemies comprise parasitoids, cleptoparasites and predators, hereafter collectively referred as "parasitoids"). Emerged individuals were determined at the species level or, if not possible, at the genus or family level. In the cases where no adult hosts emerged, hosts were identified at the genus or family level based on the features of their breeding cells (Gathmann and Tscharntke, 1999). Host abundance was defined as the total number of brood cells per habitat patch and parasitoid abundance as the number of attacked brood cells. Parasitism rate was defined as the number of attacked brood cells divided by the total amount of brood cells.

The studied trap-nesting bees and wasp species are multihabitat users that depend on scattered resources to complete their life cycle. Besides suitable food resources, they require existing cavities to build their nests. Females lay their eggs in a series of brood cells provisioned with pollen and nectar (bees) or prey (wasps). Each cell is closed with mud or organic material and the completed nest is sealed with a plug of mud or resin (Gathmann and Tscharntke, 2002; Zurbuchen et al., 2010; Coudrain et al., 2013; Bailey et al., 2014). The breeding cells are attacked by a range of enemies ("parasitoids") from several families that feed on the developing offspring or its provisions (Gathmann and Tscharntke, 1999).

STATISTICAL ANALYSES

Generalized mixed models with Poisson error distribution and site as random factor were used to test the effect of habitat amount, isolation, year (as factor with tree levels) and their pair-wise interactions on host and parasitoid abundance and species richness. To investigate if abundance and species richness of parasitoids were more affected than that of their hosts (trophic-level hypothesis, Holt et al., 1999), trophic level (parasitoids vs. hosts) and its interaction with isolation and habitat amount was included in the generalized mixed models with Poisson error distribution. Significant interactions indicate different responses to isolation and habitat amount of the two trophic groups. Generalized mixed models with binomial error distribution and patch as random factor were used to test the effect of habitat amount, patch isolation, year and their pairwise interactions on parasitism. In the models with the response variables host abundance, parasitoid abundance and parasitism, an observation-level random effect was included to account for overdispersion (Elston et al., 2001; Browne et al., 2005). The amount of habitat was square-root-transformed to improve model fit. Model fit was visually assessed by plotting fitted versus residual values and Cook's distances were calculated to detect possible outliers. Models were compared based on chi-square tests and non-significant interactions sequentially removed. To gain a more mechanistic understanding of how fragmentation drives changes in parasitoids, i.e., to test whether the fragmentation treatment effects on parasitoid richness, parasitoid abundance and parasitism were mediated by host abundance or host species richness, the best models obtained were run again with either host abundance or host species richness as covariate entered before the treatment effects in the sequentially fitted models described

above. Thus, we tested whether the variation explained by fragmentation (in the model without the co-variate host abundance or host species richness) is actually explained by the host covariate (host-mediated effects) and whether fragmentation effects still explain a significant part of the residual variation (not explained by the host covariate; direct effects not driven by host abundance or host species richness). All analyses were performed with R version 2.15.2 (R Development Core Team, 2011) using the additional package lme4 (Bates, 2012).

RESULTS

SAMPLE SIZE

A total of 41352 brood cells contained 10 bee, 28 wasp and about 50 enemy species from 21 families (**Table 1**).

EFFECTS OF HABITAT AMOUNT AND PATCH ISOLATION

Both host (Chi = 10.3, P < 0.01) and parasitoid (Chi = 21.3, P < 0.01) species richness were lower at isolated sites than at sites adjacent and connected to forest (Figures 3A,B). Host species richness was 21.4% resp. 24.3% higher at adjacent resp. connected sites compared to isolated sites, while parasitoid species richness was 68.5% resp. 64.3% higher. As predicted, the impact of isolation on species richness was significantly higher for parasitoids than for their hosts (Chi = 7.56, P = 0.02). Further, species richness of hosts (Chi = 4.6, P = 0.03) and parasitoids (Chi = 3.9, P = 0.05) increased with increasing amount of woody habitat in the surrounding landscape, without significant difference in the rate of increase between the two trophic levels (Chi = 0.07, P =0.79). There was no interactive effect of patch isolation and habitat amount on species richness of hosts (Chi = 2.1, P = 0.35) or parasitoids (Chi = 1.6, P = 0.46). Parasitoid species richness was positively correlated with abundance and species richness of their hosts (Table 2). When including host abundance or host species richness as covariate before patch isolation and habitat amount, the effects of patch isolation on parasitoid species richness

Table 1 | Insect families and overall abundance of functional groups expressed as number of brood cells summed over the three treatments "adjacent" (a), "connected" (c), and "isolated" (i) over the three years investigated.

Treatment	2008				2009			2010		
	(a)	(c)	(i)	(a)	(c)	(i)	(a)	(c)	(i)	
Host bee families	Colletidae, M	egachilidae								
Host bee abundance	196	396	374	2516	3010	2047	4709	5163	3279	
Bee parasitoid families	Anobiidae, Bombyliidae, Braconidae, Cleridae, Chrysidae, Dermestidae, Drosophilidae, Eulophidae, Gasteruptionidae, Ichneumonidae, Megachilidae, Sapygidae, Torymidae									
Bee parasitoid abundance	13	37	14	591	657	409	989	980	406	
Host wasp families	Crabronidae, Eumeninae, Pompilidae									
Host wasp abundance	1712	820	477	4240	2139	824	4066	2652	803	
Wasp parasitoid families	Bombyliidae, Braconidae, Cleridae, Chrysidae, Dermestidae, Drosophilidae, Encyrtidae, Eulophidae, Eurytomidae, Gasteruptionidae, Ichneumonidae, Pteromalidae, Sapygidae									
Wasp parasitoid abundance	302	134	56	1201	625	248	1225	670	198	



remained significant. This indicates that, in addition to indirect effects mediated by host abundance and species richness, isolation also directly affected parasitoid species richness. In contrast, habitat amount was no longer significant when host species richness was included as a covariate, which suggests an indirect effect of habitat amount on parasitoid species richness as a byproduct of the impact of habitat amount on host species richness.

Host abundance was lower at isolated compared to connected and adjacent sites (19.0% resp. 24.7% lower, Chi = 13.4, P < 0.01), but it was not related to the amount of woody habitat (Chi < 0.1, P = 0.95; **Figure 3C**). There was no significant interaction among patch isolation and habitat amount on host abundance (Chi = 1.6, P = 0.46). In contrast to host abundance, there was an interactive effect of habitat amount and patch isolation on parasitoid abundance (Chi = 6.1, P = 0.05; **Figure 3D**). Moreover, there was a strong interactive effect of patch isolation and habitat amount on parasitism (Chi = 61, P = 0.05; **Figure 3D**).

10.0, P < 0.01; **Figure 3E**). Patch isolation negatively affected parasitoid abundance and parasitism at low habitat amount, but not at high habitat amount in the landscape. This effect was robust also after accounting for variation due to host abundance (Chi = 9.2, P = 0.01).

TEMPORAL DYNAMICS

Abundance of hosts and parasitoids increased during the years of the study, mainly from the first to second year (**Table 3**). The effects of patch isolation, habitat amount and their interaction on these response variables were consistent over the three years of the study (interactions with year: all $P \ge 0.18$).

DISCUSSION

Our 3-year study demonstrates divergent effects of landscape habitat amount and patch isolation on different trophic levels of host-parasitoid communities, thereby altering parasitism rates.

Table 2 Pearson correlation coefficients between host and parasitoid
species richness and abundance and parasitism.

	Host species richness	Parasitoid abundance	Parasitoid species richness	Parasitism
Host abundance Host species richness	0.35*	0.85* 0.29*	0.66* 0.60*	0.18 0.26*
Parasitoid abundance			0.61*	0.54
Parasitoid species richness				0.33*

Significant correlations (p < 0.05) are indicated by *.

Table 3 | Results of pairwise differences between years (GLMM with Poisson or binomial distribution and site and individual observation as random effects).

	20	08–2009		2009–2010			
	Estimate	Chi	Р	Estimate	Chi	Р	
Host abundance Parasitoid	1.282 1.867	40.58 56.25	<0.01 <0.01	0.241 0.267	3.72 4.76	0.054 0.029	
Host species richness	0.356	11.51	<0.01	0.028	0.08	0.772	
Parasitoid species richness	0.461	13.25	<0.01	0.114	1.08	0.298	
Parasitism	0.678	13.02	<0.01	-0.030	0.03	0.824	

Significant results are highlighted in bold.

We found pronounced effects of patch isolation on species richness with the higher trophic level being more adversely affected than the lower one. Consistent over years, parasitism was reduced by patch isolation in landscapes with low habitat amount, but remained unaffected by isolation in habitat-rich landscapes.

Both habitat amount and isolation were important drivers of host and parasitoid species richness at a given site. Since an increase in patch isolation can be a by-product of habitat loss within the landscape (Fahrig, 2003), apparent isolation effects may simply result from lower species pool in the surrounded habitat ("sample area effect," Fahrig, 2013). However, by empirically separating habitat amount from isolation, we show that isolation per se can negatively affect species richness. In the studied host-parasitoid communities, the estimated loss of host species was 21% and that of parasitoids even 68% at isolated compared to both connected and wood-adjacent sites, challenging the view that habitat amount is the predominant driver of fragmentation effects (habitat amount hypothesis; Fahrig, 2013). As parasitoids can be both directly and indirectly (via changed host abundance or richness) affected by habitat amount and isolation, we calculated all models with and without host parameters as covariables. The significant reduction of parasitoids at isolated sites even when accounting for hosts indicates direct negative

effect of isolation on parasitoids. In contrast, the increase of parasitoid richness by habitat amount became non-significant when accounting for host richness and may thus be a purely indirect consequence of reduced host richness in landscapes with low amounts of habitat. Separate effects of landscape habitat amount and patch isolation further indicate that at least two mechanisms contributed to the species richness of the studied communities. First, landscape habitat amount should relate to the species pool that sets the number of species available to colonize a habitat patch, and second, isolation acts as an ecological filter, such that only reduced numbers of species and individuals colonize isolated habitat patches. Limited dispersal of host species, including some of the most abundant species such as the spider-hunting wasp Trypoxylon figulus (L.), which strongly declined at isolated compared to adjacent and connected sites, is probably also the main factor explaining reduced host abundance at isolated sites. Other possible explanations such as limited food resources for offspring production (e.g., Stamp, 2001) or top down pressures by parasitoids could be excluded in a recent study focusing on T. figulus as a model species (Coudrain et al., 2013). Indeed, strongly reduced parasitism rates at isolated sites would rather suggest a release of top down pressure rather than an increase (Kruess and Tscharntke, 1994; Herrmann et al., 2012; Schüepp et al., 2014b).

INTERACTIVE EFFECTS OF LANDSCAPE AMOUNT AND PATCH ISOLATION

Our results support theoretical predictions that parasitism decreases with patch isolation (Visser et al., 2009). However, we found pronounced and temporally consistent interactive effects of habitat amount and isolation on parasitoid abundance and parasitism rates. This interaction exceeds previous theoretical predictions and empirical findings. Thus the effect of patch isolation on interactions depended on the amount of habitat within the landscape. At the community level, ecological functions emerge from the interactions among multiple species (Sih et al., 1998; Kremen et al., 2007; Tylianakis et al., 2007). Therefore, changes in species richness and abundance following landscape fragmentation can induce shifts in parasitism that are unpredicted from a two species-host-parasitoid system. Moreover, the equal dispersal ability of hosts and parasitoids assumed by Visser et al. (2009) may be rather exceptional in host-parasitoid communities (e.g., Van Nouhuys and Hanski, 2002; Elzinga et al., 2007; Sivakoff et al., 2012).

Higher parasitism can result from higher parasitoid richness (Tylianakis et al., 2006; Ebeling et al., 2011; Fenoglio et al., 2012; but see Rodriguez and Hawkins, 2000). Thus, high amounts of habitat may mitigate negative isolation effect on parasitism by enhancing parasitoid species richness. Particularly when prey resources are scarce, inter-specific complementarity in prey exploitation should be maximized (Griffin et al., 2008; Tylianakis and Romo, 2010). In accordance, we observed a parallel increase in parasitoid species richness and parasitism rate with increasing amount of habitat at isolated sites where host density was low. However, parasitism decreased with the amount of woody habitat and with parasitoid richness at forest-adjacent and connected sites where host abundance was high. Thus, other drivers than parasitoid diversity must have caused the decrease of parasitism with increasing habitat amount in forest-adjacent and connected situations (see also Table 2: parasitism rate correlates with parasitoid abundance and richness, but less so with host abundance and richness). A possible explanation is that high amounts of habitat complicate host finding by the parasitoids. This may especially be true in our system in which the two most abundant host species, Osmia bicornis (L.) and T. figulus, nest in woody habitat but also forage in more open habitats (Coudrain et al., 2013). Therefore, impaired host finding success of parasitoids may release host from parasitism in complex landscapes (Gols et al., 2005; Tylianakis et al., 2007; Laliberté and Tylianakis, 2010). Additionally, a larger number of natural cavities in habitat patches surrounded by higher amount of woody habitat may have diluted parasitoid attacks on trap-nesting host individuals. Parasitoid attacks could also be reduced through better defensive strategy of aggregated hosts (Rosenheim, 1990; Rohlfs and Hoffmeister, 2004; Steffan-Dewenter and Schiele, 2008), but in our studied community, parasitism rate changed irrespective of host abundance.

Hillebrand and Cardinale (2004) provide an alternative explanation for the decrease in parasitism rate at habitat-adjacent and connected sites. They show that in contrast to parasitoid species richness, a high host species richness often implies lower parasitism rate, which is mediated e.g., by the occurrence of inedible prey or temporal inconsistency among host and parasitoid flying period. Thus, as host species richness increased with amount of woody habitat, this could explain the decrease in parasitism in connected and forest-adjacent habitats. However, the positive correlation of parasitism to host species richness lends no support to this mechanism in our study system. In summary, a positive influence of parasitoid richness at isolated sites acting in concert with impaired host finding by parasitoids in connected and forest-adjacent patches could underlie the observed modulating action of habitat amount on the relationship between parasitism and patch isolation.

IMPLICATIONS

Our study demonstrates a pronounced interactive effect between habitat amount and isolation on higher trophic level species and trophic interactions. We found that high habitat amount in a landscape can mitigate negative effects of isolation on the strength of trophic interactions. Our findings provide a possible explanation for the variable outcomes of previous studies investigating the impacts of fragmentation *per se* (e.g., Fahrig, 2003; Ewers and Didham, 2006). From a conservation perspective, a further important finding is that patches connected to small, mostly linear woody habitats showed consistently similar levels of biodiversity and ecosystem functioning (parasitism) as patches adjacent to forest. This suggests that corridors of woody habitat can effectively promote connectivity in the studied agricultural landscapes.

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