



Drivers of local densities of endangered *Lycaena helle* butterflies in a fragmented landscape

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Abstract Due to their specialised habitat requirements, butterflies are particularly vulnerable to habitat loss and fragmentation. Understanding the drivers of local abundances of species is essential for their effective conservation in fragmented landscapes. We investigated factors affecting population densities of an endangered European butterfly, the Violet Copper (*Lycaena helle*), occurring in a small metapopulation near the city of Kraków, southern Poland. The environmental parameters tested as predictors of the local densities of the species included both the variables associated with spatial structure of habitats such as patch sizes, their isolation and fragmentation as well as those potentially reflecting habitat quality. Patch area and vegetation height turned out to be the only factors significantly influencing *L. helle* densities, both having a positive effect. The positive impact of patch area is a bit surprising, since its relationship with population densities is typically negative in butterflies. In our study system it is likely to derive from source-sink dynamics as the smaller habitat patches are apparently too small to sustain viable local populations. In turn, the positive influence of vegetation height implies that the ongoing succession does not deteriorate the quality of the recently abandoned meadows yet, whereas higher turf may provide better sheltering places. The loss of almost half of *L. helle* habitat patches in the study area in recent years is alarming. However, its inclusion into the Natura 2000 system should help to conserve the species as long as this act is followed by proper management of its habitats.

Keywords Habitat fragmentation · Meadow management · Metapopulation · Source-sink dynamics · Violet Copper

Introduction

Fragmentation and habitat loss are concerned to be among the most serious threats to biodiversity (Saunders et al. 1991; Debinski and Holt 2000; Krauss et al. 2010). Butterflies belong to the organisms that suffer most from these processes, because of their relatively limited dispersal abilities and strong habitat specialization (Thomas and Hanski 1997; van Swaay and Warren 1999; Bergmann et al. 2004), which makes them an useful model group for studies on the effects of habitat loss and fragmentation on species persistence in fragmented landscapes (WallisDeVries et al. 2002). Determining factors which influence patch occupancy and local population densities are key issues in this context. Such factors can generally be divided into two main groups: (1) factors related with habitat quality (e.g., vegetation structure or resource abundance, and (2) factors reflecting spatial structure of habitat patches, mainly their area and connectivity (Thomas et al. 2001; Fleishman et al. 2002). Metapopulation theory focuses primarily on the factors belonging to the latter group. According to the theory, connectivity of habitat patches and their area are crucial predictors for the probability of species occurrence (Hanski 1999). Larger and well-connected patches are less likely to experience extinctions of local population and more likely to become recolonised, and thus more likely to be occupied, which in fact has often been confirmed empirically (Fleishman et al. 2002; Nowicki et al. 2007; Pöyry et al. 2009). In turn, habitat quality defines the carrying capacity of habitat

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patches and hence it limits local abundances of butterflies (Thomas et al. 2001; Nowicki et al. 2009).

Although the metapopulation theory in its classic form does not consider local population sizes and their densities explicitly, taking these parameters into consideration is of great importance for conservation purposes. Due to stochastic processes smaller populations are more vulnerable to extinctions (Caughley 1994; Schtickzelle et al. 2005). The risk of extinction is further increased through strong Allee effects in small populations (Cantrell and Cosner 2007). Moreover, the number of dispersers and thus the metapopulation persistence is determined not only by local population sizes, but also by their densities through density dependence of emigration rate (Nowicki and Vrabec 2011). Finally, declining population densities can typically be regarded as the first sign of deteriorating habitat quality. Detecting such negative changes and revealing the factors responsible for the process can be very helpful in planning appropriate conservation actions.

The aim of our study was to identify factors affecting local population densities of an endangered butterfly species, the Violet Copper (*Lycaena helle*). Specifically, we were interested in assessing the relative importance of factors reflecting habitat quality versus those describing spatial structure of habitat patches in shaping local abundances of the species. Due to specialised habitat requirements, the focal species usually forms metapopulations in highly fragmented semi-natural landscapes and thus it appears a good model organism for this type of research. Our ambition was also to use the results obtained for developing practical conservation recommendations.

Materials and methods

Study species

Lycaena helle is a boreo-montane species with the occurrence range extending from Central Europe, Scandinavia and Russia to the Amur Region and Mongolia (Bozano 2004; Habel et al. 2014). In Central Europe it is a post-glacial relict, inhabiting mostly highlands (Habel et al. 2011; Martin et al. 2014), although some population occupy wet lowland meadows (Skórka et al. 2007). The species is bivoltine, with adult generations occurring in spring and early summer (Settele et al. 1997; Benes et al. 2002). Its caterpillars are typically monophagous, feeding on the adderwort (*Polygonum bistorta*) in our study region (Buszko and Masłowski 2008), and thus the occurrence of the species is limited to meadows with this foodplant. Adults can also use buttercups (*Ranunculus acer* and *R. repens*) and cuckoo flowers (*Cardamine pratensis* and *C. amara*) as nectar sources (Bauerfeind et al. 2009). *Lycaena*

helle is generally regarded as a sedentary species, however it may sporadically perform long-distance movements (Habel et al. 2011). Deterioration of meadow habitats caused by land drainage and abandonment of traditional management, as well as their increasing fragmentation have been suggested as main reasons for the recent decline of the species (Fischer et al. 1999). *Lycaena helle* is currently one of the most endangered butterfly species in Europe, listed in the European Red Data Book and in the Annexes of the Habitats Directive (Van Helsdingen et al. 1996; van Swaay and Warren 1999).

Study area

The study area was located on the outskirts of Kraków, southern Poland, ca. 5–7 km southwest from the city centre (50°01'N, 19°54'E). It comprises a large wet meadow complex (ca. 800 ha) occupying the flat bed of the Vistula river valley. Its meadow habitats, including wet *Molinia* meadows (*Molinietum caeruleae* association) and lowland hay meadows (*Arrhenatheretum elatioris* association), are mostly abandoned, with a small percentage of the area being mowed annually (Nowicki et al. 2007; see this reference for a detailed description of the study area). Since 2011 a large part of the meadow complex has been protected in the Natura 2000 sites “Dębnicko-Tyniecki Obszar Łąkowy” (PLH 120065) and “Skawiński Obszar Łąkowy” (PLH 120079). Natura 2000 is a pan-European network of protected areas developed for the implementation of the key conservation policies of the European Union, namely its Birds Directive (European Commission 1979) and Habitats Directive (European Commission 1992). Each site in the network is devoted to securing a favourable conservation status of particular species or habitats endangered in Europe (European Commission 2000). Both Natura 2000 sites within our study area have been established for the conservation of four butterfly species of the Annex II of the Habitats Directive: *Maculinea teleius*, *M. nausithous*, *L. helle*, and *L. dispar* (Kudłek and Pępkowska 2008; Woyciechowski et al. 2008). Prior to the establishment of the Natura 2000 sites urbanisation was a major threat for natural values of the area (cf. Nowicki et al. 2007). Although the meadows fragments vanishing each year due to urban developments were usually small (<1 ha), they often comprised habitats of the aforementioned target species. In addition, some habitat patches of these precious butterflies were lost due to natural succession or ill-advised afforestation projects (Walasz 2008).

The study area included 21 patches with the *P. bistorta* foodplant, which can be regarded as the habitat patches of *L. helle*, when we started the butterfly surveys in 2008 (Fig. 1). Their total area encompassed nearly 23 ha, with individual patch sizes ranging between 0.08 and 2.83 ha

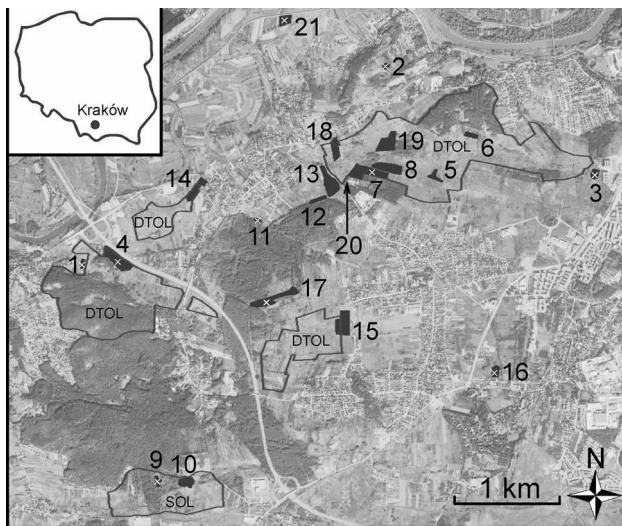


Fig. 1 Map of the investigated habitat patches of *Lycaena helle*, with the inset in the top-left corner showing their approximate location in Poland. Patch numbers correspond to those used in Table 1. The patches that ceased to exist before 2011 are marked with white crosses. *Solid lines* indicate the boundaries of separate fragments of the two Natura 2000 sites (DTOL = PLH 120065 “Dębnicko-Tyniecki Obszar Łąkowy”; SOL = PLH 120079 “Skawiński Obszar Łąkowy”)

(Table 1). The distances between the nearest neighbouring patches were typically 200 to 1500 m. Only 11 patches, though mostly the larger ones, persisted till 2011 (Table 1).

Foodplant and butterfly surveys

We mapped all the *P. bistorta* patches with ca. 1-m precision using GPS Magellan ProMark X (Magellan System Corp., USA). In order to estimate the foodplant density we counted its shoots within 3–5 randomly selected 5 × 5 m plots at each patch in June 2008 and 2011. The exact number of shoots was recorded except for a few cases in 2008, in which it was estimated with classes of ten or multiples of ten. In those cases we adopted the mid-class value when calculating the foodplant density as the mean value for both years of the study.

Butterflies surveys were conducted for the summer generation in 2008 and the spring generation in 2011, using tested and parameterized catch per-unit-time method (Nowicki et al. 2007; Niedobecki 2011). Surveys involved monitoring adult abundance and sex ratios at two representative patches through capturing individuals during 1-h capture sessions conducted in fine weather between 9:00

Table 1 Seasonal size estimates (±SE) of the investigated populations of *Lycaena helle*

Patch number	Patch area (ha)	Summer generation 2008	Spring generation 2011
1	0.39	0 ± 0	–
2	0.08	20 ± 8	–
3	0.42	42 ± 7	–
4	2.30	219 ± 44	–
5	0.42	77 ± 8	28 ± 3
6	0.36	52 ± 6	28 ± 1
7	2.67	354 ± 113	–
8	1.59	1999 ± 320	135 ± 8
9	0.56	63 ± 6	–
10	1.11	139 ± 21	105 ± 21
11	0.11	13 ± 2	–
12	0.52	291 ± 6	37 ± 4
13	2.77	262 ± 24	306 ± 27
14	0.62	74 ± 20	44 ± 3
15	2.83	976 ± 577	286 ± 6
16	0.20	19 ± 3	–
17	1.72	210 ± 88	–
18	0.86	204 ± 6	81 ± 23
19	2.07	321 ± 48	190 ± 59
20	0.26	32 ± 4	31 ± 2
21	0.69	0 ± 0	–
Total	22.53	5367 ± 680	1271 ± 73

and 17:00. The capture sessions were held daily to every second day throughout the entire flight period (27 June to 3 August 2008, and 8 May to 9 June 2011). Peak occurrence periods of adults in both seasons (summer 2008 and spring 2011) were determined on the basis of the highest daily numbers of captured individuals and their balanced sex ratio, i.e., approximately 60 % proportion of males concerning their higher catchability (Nowicki et al. 2005a). Subsequently, during these peak adult occurrence periods we conducted 1-h capture sessions at each habitat patch for 2 days (8–9 July 2008, as well as 24 May and 26 May 2011). While the data collected in this way were not comprehensive enough to yield seasonal population sizes with mark-capture models, they allowed the estimation of peak population sizes based on capture frequencies (cf. Nowicki et al. 2007). Peak population size estimates were subsequently extrapolated into seasonal population size estimates following the approach described by Nowicki et al. (2005a), which accounts for flight period length and average adult life span. The latter was calculated as $e = (1 - \phi)^{-1} - 0.5$ (Nowicki et al. 2005b), where ϕ represents mean daily survival rate derived with the Cormack–Jolly–Seber mark-recapture model (Schwarz and Arnason 1996; Schwarz and Seber 1999) using the MARK 5.1 software (White and Burnham 1999). The adult life span estimates reached 4.99 and 5.93 days in summer 2008 and spring 2011 respectively.

Statistical analysis

The population size estimates were converted into seasonal densities through division by patch area. In several

temperate zone butterflies spring generations were found to be several times smaller than summer ones (Fric et al. 2006; Pickens 2007), which was also the case in our study on *L. helle*. Thus, in order to make possible the comparison of seasonal densities recorded at particular patches in summer 2008 and spring 2011, we used their standardised values. Standardisation was done through dividing densities recorded at each patch by mean density across all the patches in a given season. Subsequently, for each patch we calculated the mean of the standardised density values obtained for summer 2008 and spring 2011. In this way we derived relative density index, which was used as the dependent variable in the analysis of factors affecting local abundances of *L. helle*. For the patches that ceased to exist before 2011, i.e., in the cases with no standardised density value available for spring 2011, the standardised density values obtained for summer 2008 were adopted as the relative density index.

We opted for deriving a combined density index for both investigated seasons instead of analysing the patterns in local abundances of the focal species separately for summer 2008 and spring 2011, because the population sizes recorded at each habitat patch within a 3-year year period are likely to be non-independent. In other words, we wanted to avoid the problem of temporal autocorrelation in local population densities, which should be expected within such a short period (cf. Royama 1992; Liebhold et al. 2004), even though it could not be assessed with the data from just two seasons. In turn, it must be underlined that there was absolutely no indication of spatial autocorrelation in local abundances of *L. helle* across the investigated patches. Mantel test r values were very close to zero for

Table 2 Habitat patch parameters used as predictors in the analysis of local densities of *Lycaena helle*

Parameter	Description	Interpretation
Parameters describing spatial structure		
Patch area	Area covered with the <i>Polygonum bistorta</i> foodplant (ha)	Measure of habitat patch size
Connectivity	Hanski et al. (1994) connectivity index I_3 defined as $\sum \exp(-d_{ij})$ where d_{ij} is the distance (km) of patch i to other patches ($i \neq j$)	Measure of patch isolation; decreases with increasing isolation
Compactness	Mean distance of points within patch from its edge divided by the square root of patch area	Measure of patch fragmentation; decreases with increasing fragmentation
Parameters describing habitat quality		
Foodplant abundance*	Density of <i>Polygonum bistorta</i> shoots (per m ²)	Measure of foodplant availability
Vegetation height	Mean height of meadow vegetation within patch (cm)	Measure of meadow succession
Distance to stream/ditch*	Mean distance of points within patch to the nearest stream or ditch (m)	Measure of habitat moisture
Distance to building/road	Mean distance of points within patch to the nearest building or road (m)	Measure of potential anthropogenic pressure

* The parameters that were logarithmically transformed in order to achieve their normal distributions

population densities recorded in both seasons (summer 2008: $r = 0.0729$, $P = 0.2035$; spring 2011: $r = -0.0439$, $P = 0.3471$) as well as for the combined density index ($r = 0.0631$, $P = 0.2557$).

The factors tested as predictors of local abundances of *L. helle* included both the parameters associated with spatial structure of habitats such as patch sizes, their isolation and fragmentation as well as those potentially reflecting habitat quality, including anthropogenic pressure (see Table 2 for details). Except for the foodplant density and vegetation height, the predictor values were derived from the existing GIS database of the study area, using the Idrisi 2.0 software (Eastman 1997). Foodplant density and distance to streams were logarithmically transformed in order to achieve their normal distributions. When estimating the connectivity of each habitat patch, which depends on the distances to all other patches in the system (Hanski et al. 1994), we used the spatial configuration of the patches existing in summer 2008. However, with this parameter based on the patch configuration in spring 2011, which we also attempted for comparison, the outcome of all the analyses remained virtually unchanged.

All the parameters used as predictors were only moderately correlated with one another (Pearson's r invariably below 0.5, thus within the range in which variables in multivariate analyses may be regarded as fairly independent; Legendre and Legendre 1998). Their effect on local abundances of *L. helle* was analysed with multiple linear regression model with a forward stepwise variable selection. Nevertheless, it should be stressed that a backward elimination procedure would not make any difference as it yielded identical final results. The statistical analyses were performed using the Statistica 10 program (Statsoft 2010), except for the Mantel tests, which were conducted with the help of the program Zt (Bonnet and Van de Peer 2002).

Results

In summer 2008 *L. helle* was found in 19 out of 21 habitat patches, while in spring 2011 it occupied the 11 remaining patches (Table 1). During the capture sessions we recorded 938 adults in summer 2008 and 241 adults in spring 2011. The total metapopulation size was estimated at 5367 adult

butterflies for the summer 2008 generation, with ten relatively large local populations (>100 adults) encompassing together 93 % of all individuals. Seven of these large populations persisted till spring 2011, when they jointly constituted 90 % of the total metapopulation estimated at 1271 individuals. Butterfly densities at particular patches in summer 2008 varied greatly between 95 and 1259 individuals per ha (coefficient of variation $CV = 0.91$). In spring 2011 they were much more uniform, ranging between 67 and 119 individuals per ha ($CV = 0.19$).

The multiple regression analyses, using both stepwise and backward variable selection, indicated that local densities of *L. helle* were significantly influenced by patch area and vegetation height, and the density index positively correlated with both above variables (Table 3). The results of the stepwise variable selection procedure showed that patch area explained 24.4 % of the variation in the density index, and vegetation height explained further 20.2 % (Table 3). The overall model fit was thus moderately good, reaching 44.6 %.

Discussion

The outcome of our study allows a reliable assessment of the status of *L. helle* metapopulation within the study area. From the conservation point of view our results are of serious concern. Obviously, the fact that the metapopulation size estimated in summer 2008 was approximately four times larger than the one in spring 2011 should not necessarily be perceived as an indication of a sharp decline in *L. helle* abundance within the investigated period. Instead, it is likely to reflect the difference between summer and spring generations, as the former ones are typically several times larger in bivoltine butterflies of the temperate zone (Fric et al. 2006; Pickens 2007). However, disappearing of more than 40 % of occupied patches within the 3 years of our study, even though they were generally of low quality and inhabited by small populations, is alarming. These 'lost' patches encompassed 36 % of area occupied by *L. helle* and 17.5 % of its seasonal numbers in 2008.

The fact that patch area explained a considerable fraction of variation in *L. helle* densities across the habitat

Table 3 Results of multiple regression analysis of factors affecting local densities of *Lycaena helle*

Parameter	Slope (\pm SE)	Intercept (\pm SE)	t	P	Adjusted R^2
Constant		-0.124 ± 0.252	-0.489	0.6316	
Patch area	0.809 ± 0.208	0.301 ± 0.078	3.880	0.0015	0.244
Vegetation height	0.545 ± 0.217	0.004 ± 0.002	2.514	0.0195	0.202
Total					0.446

Only significant factors are included

patches seems to be a bit surprising, mainly because of the positive relationship. While patch area is a good predictor of both the probability of species occurrence and population size (Hanski et al. 1995; Hill et al. 1996; Bauerfeind et al. 2009), it rarely correlates with population density. Moreover, the result of the meta-analysis by Hambäck and Englund (2005) suggests that whenever such a correlation exists in butterflies it is generally negative. A possible explanation for the positive correlation between patch area and *L. helle* densities that we recorded may be a situation, in which *L. helle* in our study area does not exist in a metapopulation structure but rather in a source-sink system (sensu Pulliam and Danielson 1991).

Obviously, the data we have gathered do not allow proving the existence of such a system, as it would require profound information about population dynamics within habitat patches and dispersal among them. Nevertheless, some indirect evidence makes plausible our hypothesis concerning possible source-sink dynamics in the investigated system. First of all, the small habitat patches may actually be too small to sustain *L. helle* populations, and thus individuals observed at such patches may mostly represent immigrants from larger patches (cf. Brückmann et al. 2010). This postulate is supported by the fact that spring population sizes in several small patches reached only ca. 30–40 individuals. Due to temporal fragmentation of the populations, i.e., the fact that only a fraction of individual is present at any day of the season as their life span is much shorter than flight period length (Nowicki et al. 2005b), such low numbers make finding a mating partner difficult (and sometimes even impossible), thus negatively affecting population growth.

The positive relationship between patch area and *L. helle* densities is also likely to be supported by asymmetric dispersal. A positive effect of patch area on immigration probability and its negative effect on emigration rate have often been reported for butterflies (Wahlberg et al. 2002; Rabasa et al. 2008; Bonelli et al. 2013). Consequently, large patches should mostly gain dispersing individuals, whereas the small ones should mostly lose them. However, our argument that the observed positive density-area relationship may derive from asymmetric dispersal can be valid only if there is considerable dispersal among the local populations. Such an assumption seems to contradict the strong site-fidelity reported for *L. helle* (Habel et al. 2011). However, our anecdotal observations of inter-patch movements over a few hundred meter distances (Nabielec and Nowicki, unpublished data) as well as the results of a recent quantitative analysis of the species mobility (Craioveanu et al. 2014) indicate that *L. helle* may be more mobile than it is popularly believed.

A positive correlation between meadow vegetation height and *L. helle* densities was also detected. Negative

relationship would indicate that succession may threaten the persistence of *L. helle* populations. Nevertheless, an earlier study in the same region showed that in its initial stages the succession at recently abandoned wet meadows improves habitat quality for many butterflies, although in later stages its effect is definitely detrimental (Skórka et al. 2007). Additionally, higher vegetation can act as shelter from bad weather conditions e.g., strong wind (Dennis 2004), and specifically in *L. helle* it can be used for roosting at night or in poor weather (Turlure et al. 2009; Goffart et al. 2014).

The lack of the effect of foodplant availability on *L. helle* densities may seem a bit surprising, but it most likely reflects the fact that *P. bistorta* is overabundant at the investigated patches (reaching 3–263 shoots per m²) and it does not limit the butterfly populations. The species densities were also not influenced by patch connectivity, but again connectivity does not appear to be a limiting factor in our study system, because most of the habitat patches (especially the centrally located large ones) are rather well-connected. However, it must be noted that studies focusing on *L. helle* patch occupancy indicated positive effects of patch sizes and their connectivity (Fischer et al. 1999; Bauerfeind et al. 2009).

To conclude, local densities of *L. helle* proved to be influenced by both spatial structure of habitat patches, as indicated by a significant effect of patch area, as well as by habitat quality, represented by vegetation height. Such an outcome is in contrast with the findings of our earlier study on the abundance patterns on *Maculinea* butterflies, for which only spatial structure of habitat patches turned out to matter (Nowicki et al. 2007). The implication is that in order to support the persistence of *L. helle* within the investigated region the conservation efforts are needed both at landscape scale, so as to ensure the existence of a number of large habitat patches, as well as at within-patch scale, where they should be focused on improving habitat quality.

The inclusion of the majority of studied area into the Natura 2000 network offers a chance for improving the status of *L. helle* in the region. Thus, from the conservation point of view, the priority should be preserving the group of large and well-connected patches of *P. bistorta* in the central part of the meadow complex (see Fig. 1). The remaining patch (number 10 in Fig. 1), although relatively large as well, is probably too isolated to effectively serve as a source of immigrants to other populations. According to Hanski et al. (1996) a stable metapopulation should ideally consist of at least 15 well-connected populations, while the aforementioned key group of patches hosts only ten populations. Therefore it is vital that every single of them is preserved, including the two (patches 12 and 13 in Fig. 1), which are located directly outside the boundary of the

Natura 2000 site “Dębnicko-Tyniecki Obszar Łąkowy”. This should serve as a strong argument for extending this Natura 2000 site so as to include the focal meadow fragment, which in fact was the case in the initial proposal for the site (Kudłek and Pępkowska 2008).

Obviously, the inclusion into the Natura 2000 network will help *L. helle* population only if it is followed by proper management of their patches. Rotational mowing and extensive grazing are believed to be the most effective management methods for *L. helle* habitats (Fischer et al. 1999; Bauerfeind et al. 2009). Incorporating such management into action plans for the meadows occupied by *L. helle* will surely contribute to its more effective conservation. Moreover, these types of land management are also favourable for many other butterfly species (Elligsen et al. 1997; Meyer-Hozak 2000; Dover et al. 2011).

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