



## Adjacent woodlands rather than habitat connectivity influence grassland plant, carabid and bird assemblages in farmland landscapes

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6   Adjacent woodlands rather than habitat connectivity influence grassland plant, carabid and bird  
7   assemblages in farmland landscapes

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## **Abstract**

One response to biodiversity decline is the definition of ecological networks that extend beyond protected areas and promote connectivity in human-dominated landscapes. In farmland, landscape ecological research has focused more on wooded than open habitat networks. In our study, we assessed the influence of permanent grassland connectivity, described by grassland amount and spatial configuration, on grassland biodiversity. We selected permanent grasslands in livestock farming areas of north-western France, which were sampled for plants, carabids and birds. At two spatial scales we tested the effects of amount and configuration of grasslands, wooded habitats and crops on richness and abundance of total assemblages and species ecological groups. Grassland connectivity had no significant effects on total richness or abundance of any taxonomic group, regardless of habitat affinity or dispersal ability. The amount of wooded habitat and length of wooded edges at the 200 m scale positively influenced forest and generalist animal groups as well as grassland plant species, in particular animal-dispersed species. However, for animal groups such as open habitat carabids or farmland bird specialists, the same wooded habitats negatively influenced richness and abundance at the 500 m scale. The scale and direction of biodiversity responses to landscape context were therefore similar among taxonomic groups, but opposite for habitat affinity groups. We conclude that while grassland connectivity is unlikely to contribute positively to biodiversity, increasing or maintaining wooded elements near grasslands would be a worthwhile conservation goal. However, the requirements of open farmland animal species groups must be considered, for which such action may be deleterious.

## **Key words**

Multi-taxon biodiversity; habitat network; random forests; multi-model inference; dispersal; hedgerow; human-dominated landscapes

45    **Highlights**

46    Permanent grassland connectivity had little effect on biodiversity

47    Wooded habitats in the landscape influenced grassland plants, carabids and birds

48    Open habitat animal specialists avoided landscapes with woods and hedgerows

49    Response to landscape was more dependent on habitat affinity than on taxonomic group

50    Increasing or maintaining wooded elements near grasslands may be good policy

51

## 1. Introduction

In human-dominated, highly fragmented landscapes, site protection alone is unlikely to achieve the objective of stopping or reversing biodiversity declines (Bennett and Saunders 2010). Rather enhancing habitat connectivity or linkages, such as ecological corridors and networks, will be needed to ensure the viability of plant and animal populations (Crooks and Sanjayan 2006). This connectivity depends on the amount (area) and spatial configuration of suitable habitat as well as on matrix permeability (Bennett 2003). Nature conservation policy has accordingly evolved towards the implementation of habitat networks (ecological networks or greenveining), which are increasingly integrated into international, national or regional planning; particularly in Europe (Boitani et al. 2007; Jongman et al. 2011). For example, since 2009 in France, it has become a legal requirement that habitat networks be established at national, regional and local levels (Vanpeene-Bruhier and Amsallem, 2014). Most territory may potentially be concerned by this policy's implementation, including areas subject to more or less intense human activity. The need to develop nature conservation outside protected areas has been accompanied by the recognition that most conservation efforts have tended to focus on rare species and habitats, sometimes to the detriment of common biodiversity (Inger et al. 2015). However, such biodiversity dominates ecosystem composition and makes an important contribution to ecosystem functioning (Gaston and Fuller 2008; Gaston 2008).

Agriculture is clearly one of the major human activities to be concerned by these shifts in policy. Farming landscapes occupy large areas of land (approximately 75% in Europe, Robinson and Sutherland 2002, and up to 40% globally, Foley et al. 2005) and are associated with a valuable and partly typical flora and fauna (Benton et al. 2003), which has declined severely over the past few decades (Chamberlain et al. 2000; Donald et al. 2001; Liira et al. 2008). There is some consensus on the importance of landscape complexity, or the amount of semi-natural habitat, in determining levels of biodiversity in farmland landscapes (Benton et al. 2003; Tschardt et al. 2012). In this context, attempts are being made to preserve and enhance semi-natural habitats of ecological value, like hedgerows and permanent grasslands. These habitats have generally diminished (Meeus 1993) and are

the focus of some considerable attention in the context of nature conservation in farming areas. For instance, in France permanent grasslands declared by farmers represented 6.7 million ha of agricultural land at national scale in 2006. By 2010 6.3% of them had been lost, replaced by temporary grasslands or scrubland (Faïq et al. 2013). Enhanced connectivity of these semi-natural habitats is expected to counteract the negative effects of such reductions in biodiversity (e.g. Hendrickx et al. 2007).

Studies of connectivity effects in farmland contexts have tended to focus on wooded habitat types (e.g. Gil-Tena et al. 2014; Neumann et al. 2016). The effectiveness of linear woodland habitats, like hedgerows, as corridors facilitating forest specialist dispersal has been demonstrated for a diversity of taxonomic groups: birds (Davies and Pullin 2007), arthropods (e.g. carabid beetles, Petit 1994; Millan-Pena et al. 2003), plants (Jamoneau et al. 2011) and small mammals (Gelling et al. 2007). Meanwhile the possible effects of increasing permanent grassland connectivity for farmland biodiversity have received relatively less attention. Most studies of grassland connectivity have concentrated on rarer grasslands of high conservation value, such as wet grasslands (e.g. Lafage et al. 2015), calcareous grasslands (e.g. Brückmann et al. 2010; Rösch et al. 2013), heathland (e.g. Piessens et al. 2005), species-rich grasslands at higher altitudes (e.g. Marini et al. 2008) or indigenous grasslands fragmented by commercial forestry (e.g. Samways & Pryke, 2016). These studies, involving a diversity of taxonomic groups, have shown that species of open, grassland habitats may respond positively to increasing grassland connectivity. Common permanent grasslands of agricultural mosaics, which are mesophilic, managed grasslands that have not been ploughed for several years, have been understudied in landscape connectivity research. With their lack of vertical heterogeneity and dependence on highly variable forms of management, such grasslands are difficult to describe (Allen et al. 2011) and until the recent focus on common biodiversity may have been expected to harbour few species of conservation interest. Only a few studies in Europe have begun to explore how connectivity of more managed grasslands may influence arthropods and plant assemblages (Öckinger et al. 2012; Wamser et al. 2012; Villemey et al. 2015). These studies found contrasting effects of grassland connectivity on species richness. Thus, it is important to further explore to what extent

connectivity of common permanent grassland contributes to maintaining biodiversity in farmland mosaics.

The effects of landscape context depend on the varying scales of perception of taxa with differing dispersal abilities (Jackson and Fahrig 2012) and within-taxa differences in dispersal ability may potentially obscure the effects of connectivity at assemblage level (e.g. Wamser et al. 2012). In addition, species with different habitat affinities may display opposing responses to landscape structure (Mauremooto et al. 1995; Thomas et al. 1998; Filippi-Codaccioni et al. 2010). Therefore, using two spatial scales, we aimed to assess how variation in grassland connectivity (amount and spatial configuration) influenced the richness and abundance of various taxonomic groups with potentially different scales of landscape perception: plants, carabids and birds observed in common, mesophilic, permanent grasslands. We hypothesized that grassland connectivity would increase richness and / or abundance of these assemblages. We also looked at the richness and abundance of species groups based on habitat affinity and dispersal ability, and compared the influence of grasslands with that of wooded habitats and crops in the surrounding landscape. We further hypothesized that grassland specialist or open habitat species would benefit most from grassland connectivity or amount of crops. Wooded habitats (particularly hedgerows) may act as barriers for these species, while they may help forest species to disperse and therefore increase richness and abundance of this group in grasslands. We expected a stronger influence of landscape structure on species with low dispersal ability.

## **2. Method**

### **2.1. Study areas and permanent grassland selection**

The study was conducted in north-western France, where farmland landscapes contain annual crops (mostly winter cereals, but also spring maize and oilseed rape), along with temporary and permanent grasslands (not ploughed for at least 5 years), separated by woodland and hedgerow networks (which will be referred to as wooded habitats). We selected four study areas located close to Nantes, Angers,

La-Roche-sur-Yon and Rennes which shared the same climate (temperate oceanic), geo-morphological conditions (acidic substrate: schists, granites, sandstones) and a similar agricultural history of mixed dairy farming systems (Appendix 1A). In these areas, we selected permanent grasslands to maximize variation in the proportion of permanent grassland in the surrounding landscape. These production grasslands were originally established by farmers, using a mixture of sown species of grasses and sometimes clover, but most plant species were native. In western Europe and elsewhere, such grasslands have a long history of agricultural management, and their biodiversity has long been determined by the dynamics of regular mowing and grazing. To minimise other sources of variation, we excluded wet grassland and grasslands that were close to major roads or urban infrastructure. All were grazed, as this factor is known to have a strong influence on grassland plant assemblages (Gaujour et al., 2012).

## 2.2. Sampling and biodiversity measures

Biodiversity was sampled in between 21 and 55 permanent grasslands, depending on taxonomic group. The proportion of grassland in the landscape surrounding sampled grasslands (200m-radius buffer) ranged from 12 to 85% for plant, from 10 to 58% for carabids and 28 to 85% for birds. Plants were sampled in 55 permanent grasslands (4 Nantes, 18 Angers, 14 La Roche-sur-Yon, 19 Rennes). Each grassland was visited once between 2011 and 2015, in the late spring or summer (May –July). All vascular species were listed within 3 quadrats of 2 x 2m per grassland, placed at least 2 m from the field edge, and these were subsequently pooled for calculation of species richness measures.

Carabid beetles were sampled in 40 permanent grasslands (40 Rennes), using pitfall traps. Two pitfall traps per grassland were located 1m from each other and 10m from grassland edge. Carabids were sampled during six sampling periods, two per month in May, June and September 2011, to coincide with the seasons during which carabid beetles emerge (spring and late summer). Traps were collected every two weeks, after being open for seven consecutive days. Eliminating a few lost traps (destroyed by cattle), we measured activity-density as the number of individuals per trap per week and we



checked that the number of valid traps did not influence species richness. The six sampling periods were then pooled to calculate species richness and activity-density measures for each grassland.

Birds were sampled in 21 permanent grasslands (10 Angers, 11 La Roche-sur-Yon) and their associated field margin vegetation, using standard territory mapping (Bibby et al. 2000). The grasslands were visited six times per breeding season in 2014 and 2015 (between mid-March and mid-June). All bird surveys were carried out by a single observer between 1 and 4 hours after sunrise, on days without continuous rain or wind. Species defending one or more territories in at least one of the two years were considered to be breeding birds. The two years were pooled for calculation of species richness measures while abundance values corresponded to the maximum number of territories a species defended over the two years.

Plants, carabids and birds were classified according to their habitat affinity (generalists, grassland or open farmland specialists, woodland specialists) or to their dispersal ability (dispersal mode for plants, wing system for carabids and a morphological dispersal ability predictor for birds; see below). Species richness and abundance (for carabids and birds) were calculated for each habitat affinity or dispersal ability group. Plant ecological preferences were adapted from Baseflor database (Julve, 1998) and plant dispersal mode extracted from Bioflor (Kuhn et al. 2004) and Baseflor (Julve 1998) databases (Appendix 1B). Carabid habitat affinity data were adapted from Neumann et al. (2016) and Roger et al. (2010). Carabid species dispersal ability was estimated by the wing system (Kotze and O'Hara 2003; Purtauf et al. 2004) using information from Barbaro and van Halder (2009), Ribera et al. (2001), and BETSI database (2012, Appendix 1C). Bird habitat affinity groups were based on analysis of national bird monitoring data (Jiguet 2010) except for 14 missing species, for which we used regional atlas descriptions of habitat use (Marchadour 2014). Bird dispersal ability was estimated by using the quotient of Kipp's distance (distance between the tip of the first primary feather to the tip of the wing) and bill depth (measured at the proximate edge of the nostrils), which has been shown to be a good surrogate for natal dispersal distance in European passerines (Dawideit et al. 2009). Biometric data were provided by the Senckenberg Biodiversity and Climate Research Centre in Frankfurt, Germany

(for details, see Laube et al. 2013). Based on the obtained dispersal ability predictor we classified birds into 4 dispersal categories (low, medium, high and long distance) using Jenks natural breaks (Appendix 1D). Some bird ecological groups containing very few species, at low levels of frequency or abundance, were excluded from analysis (urban species and long-distance dispersers).

### 2.3. Land-cover maps and landscape descriptors

We produced land-cover maps of the four study areas using existing land cover databases coupled with automatic classification of satellite imagery, and corrected by photo-interpretation and ground-truthing. Land-cover classification was as follows: built-up area (roads and buildings), water bodies, crop fields, permanent grasslands, and wooded habitats (woodland and hedgerows). We paid particular attention to the accurate estimation of permanent grassland distribution, using photo-interpretation and the Land Parcel Identification System (i.e. Registre Parcellaire Graphique), where farmers subsidised by the Common Agricultural Policy (CAP) declare field use. These maps were produced using QGIS (QGIS Development Team, 2015).

Landscape descriptors were calculated from these maps within 200m and 500m radius, circular buffers centred on each sampled grassland, in order to describe two components of the connectivity of permanent grassland, i.e. grassland amount and grassland configuration. These circular buffers were centred on field centroids for plants and birds, or on exact trap locations for carabids (recorded with a GPS). The choice of two spatial scales was to account for expected differing grains of landscape perception between plants, carabids and birds (Jackson and Fahrig, 2012). Amount of permanent grassland was described as the proportion of surface area covered by these habitats in buffers.

Configuration of permanent grassland was assessed via three contrasting landscape descriptors, chosen to investigate the possible influence of different forms of spatial arrangement: largest grassland patch area, length of permanent grassland edges and a grassland connectivity index (derived from Hanski 1999, see Steffan-Dewenter 2003):

$$C_k = \sum_i^n e^{-D_{ik}} A_i$$

where  $k$  is the focal sampled permanent grassland,  $n$  the number of other (non-sampled) permanent grasslands in the 200 m or 500 m scale landscape,  $D_{ik}$  is the distance between the sampled grasslands  $k$  and the neighbouring permanent grassland  $i$  and  $A_i$  the area of the neighbouring permanent grassland  $i$ . This measure of connectivity increases when many, large, permanent grasslands are located near the sampled grassland. The proportion of other main habitat types was also calculated for the two buffer sizes: wooded habitat (woodlands + hedgerows), and crops (including temporary grasslands which are part of crop rotations). Total length of wooded habitat edges was also measured as it reflects fragmentation of open habitats and adjacencies between open and wooded habitats. In the studied landscapes, wooded habitat edges mostly corresponded to the presence of hedgerows. Landscape descriptors were calculated at the two spatial scales using Chloé 2012 (Boussard and Baudry 2014). The extent of variation in each landscape descriptor, obtained at each scale and for each study taxon, is presented in Appendices 1E and 1F.

#### 2.4. Statistical analyses

To test the representativeness of our sampled communities compared to potential richness, we calculated non-parametric species richness estimators of Chao2 across all grasslands (Chao, 1987). These estimated richness values were compared to the observed total number of species, for each taxon. We tested the effect of landscape descriptors on biodiversity measures using two successive steps (e.g. see Puech et al. 2014, Aviron et al. 2016): (i) preselection of landscape descriptors with random forest procedure, (ii) multi-model inference (MMI) and averaging of multiple regression models. Pearson's correlations between landscape descriptors and across scales are presented in Appendices 1G-I.

All seven landscape descriptors at both scales (14 variables) were included in a random forest analysis (Breiman 2001; Strobl et al. 2009), a recursive partitioning method recommended to deal with “small n large p problems” (i.e. few replicates and many environmental variables), complex interactions and correlated environmental variables (Strobl et al. 2008). For each biodiversity measure, 10,000 trees were grown and landscape descriptor importance was evaluated as the difference in model accuracy

before and after 10 permutations of values of the considered descriptor, averaged over all trees. Conditional importance that adjusts for correlations between environmental variables was used. The absolute importance value of the lowest negative-scoring landscape descriptor was used as a threshold to determine relevant and informative variables to retain for regression models (for full details, see Strobl et al. 2008, 2009). Landscape descriptors selected for each biodiversity measure may be found in Appendices 1J-L. To assess the influence of landscape descriptors on the global assemblage, the conditional importance values of each variable were averaged across the biodiversity measures for each taxon.

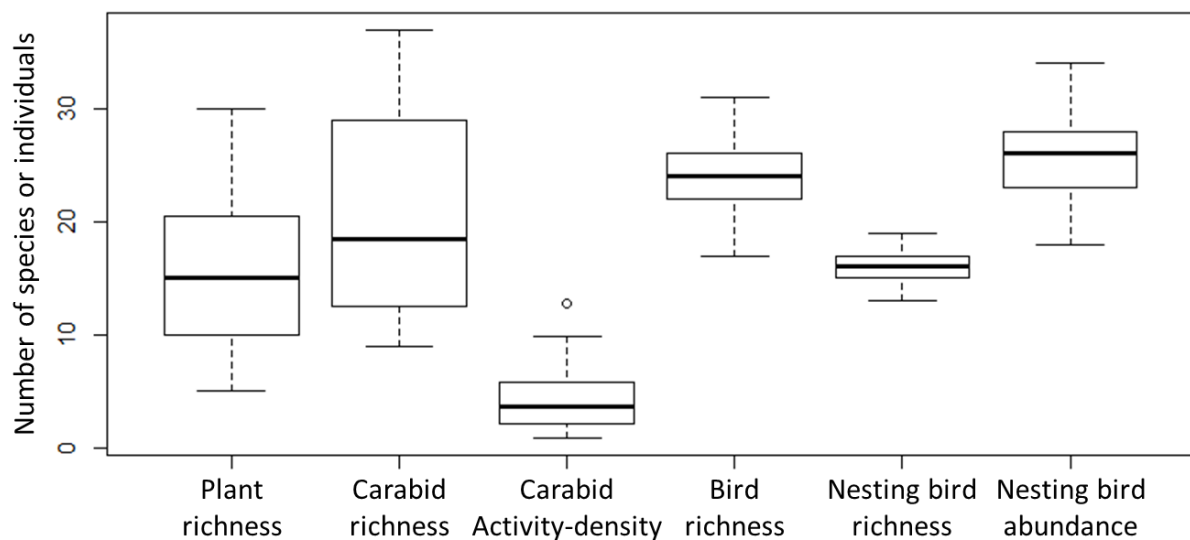
The selected landscape descriptors were then included in multiple regression models, which was analysed using multi-model inference (MMI) and model averaging. MMI analysis deals with model selection uncertainty (Burnham and Anderson 2002; Arnold 2010) and is robust against correlation among descriptors (Smith et al. 2009, 2011). All landscape descriptors were mean-centred and divided by the standard deviation to make the coefficients comparable (Smith et al. 2009, 2011). Following the MMI procedure, we created linear models for each possible combination of landscape descriptors and ranked them based on the corrected Akaike information criterion (AICc). Then, we computed standardised average regression coefficients weighted by the Akaike weights across supported best models ( $\Delta AICc < 4$ ) and tested their significance using unconditional 95 % confidence intervals (Burnham and Anderson 2002; Smith et al. 2009, 2011). Averaged model coefficients and their 95 % confidence intervals are presented in Appendices 1M-P. We also checked that the detected effects were consistent across all individual models included in model averaging (appendix 2). We more specifically looked at the correct estimation of relative effects of correlated landscape descriptors (grassland-related vs. wooded habitat descriptors, composition vs. configuration descriptors, and between scales).

Residuals of averaged models were tested for normality (Shapiro-Wilcoxon test and quantile-quantile plots) and spatial auto-correlation (Moran correlogram). When residuals were not normally distributed a new average model was built using the adequate distribution using a generalized linear model, either

Poisson distribution for non-overdispersed data or negative binomial distribution for overdispersed data (Crawley 2007; Bouche et al. 2009). No model showed spatial auto-correlation. Models included study area (plants and birds) or locality (carabids) as random factors (mixed models). All statistical tests were performed using R software 3.3.1 (R Core Team 2016) using the ‘vegan’ package for Chao2 estimation (Oksanen et al. 2013), the ‘party’ package for random forest analyses (Hothorn et al. 2013), the ‘MuMin’ package for MMI analyses (Barton 2016), and the ‘lme4’ package for generalized linear mixed-effects models (Bates et al. 2015), the ‘qcc’ package for over-dispersion testing (Scrucca 2004), and the ‘ncf’ package for spatial-autocorrelation test (Bjornstad 2016).

### 3. Results

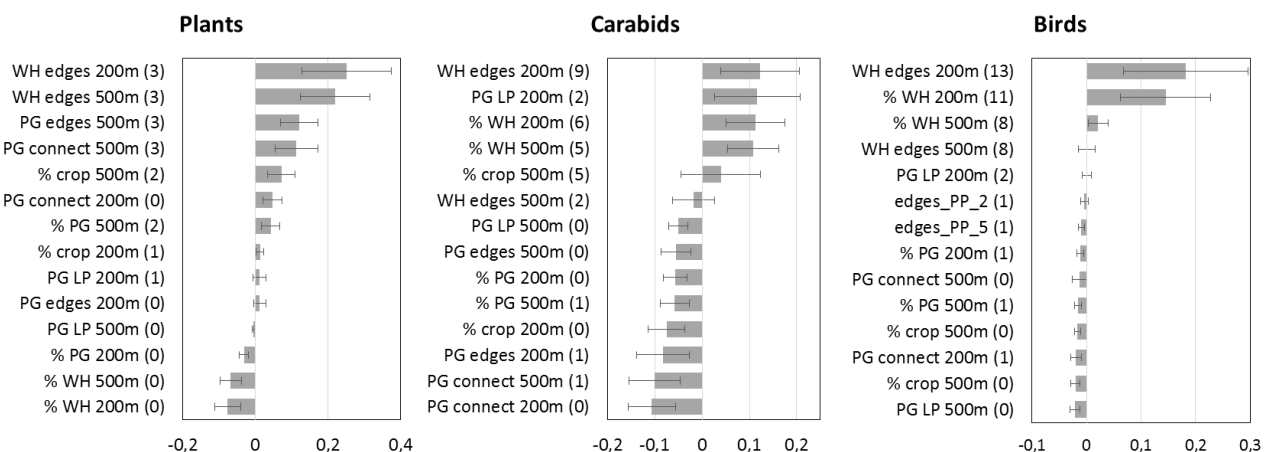
In total, we sampled 108 plant species ( $\text{Chao2} = 125.0 \pm 8.2$ ), 76 carabid beetle species ( $\text{Chao2} = 84.3 \pm 5.5$ , 1922 individuals) and 63 bird species ( $\text{Chao2} = 73.4 \pm 7.3$ ), 41 of which were breeding ( $\text{Chao2} = 54.7 \pm 10.7$ , 672 breeding territories). Observed species richness was therefore close to expected for all taxonomic groups, indicating that the sampling intensity was adequate.



**Figure 1.** Boxplot of plant, carabid and bird biodiversity measures per sampled permanent grassland, all species included.

Measures of richness and abundance at the assemblage level varied between sampled permanent grasslands (Fig. 1), but could not be explained by permanent grassland connectivity. Proportion and spatial configuration descriptors of permanent grassland were sometimes selected by the random forest

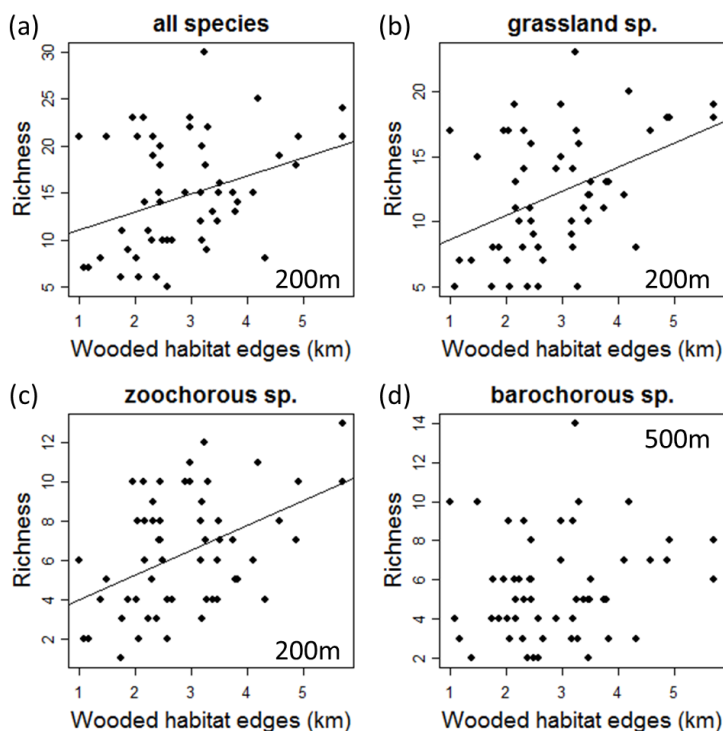
analysis, at both spatial scales, but they had limited explanatory power for carabid and bird assemblages (Fig. 2). The only exception was largest grassland patch area at the 200m scale for carabids, but this descriptor was selected only twice (Fig. 2). For plants, grassland-related descriptors were of higher importance than for the animal groups and were often selected, particularly the configuration descriptors at the 500m scale (Fig. 2). However, they never significantly affected total species richness or activity-density / abundance of plants, carabids or birds (Appendices 1M-P). Nor did they have any significant influence on species richness or abundance measures of the different ecological groups, based on habitat affinity or dispersal ability. The only exception was a nearly significant negative effect of largest grassland patch area on total richness of forest bird specialists, at the 200m scale (Appendix 1O). So, despite considerable variation in the amount and configuration of permanent grasslands across sampled grasslands (e.g. 10-80 % of permanent grassland for plants at the 200 m scale, Appendix 1E), these factors had little or no influence on the different taxonomic groups.



**Figure 2.** Conditional importance scores of landscape descriptors provided by random forest analyses, averaged across biodiversity measures for each studied taxon. Error bars are the standard errors.  $n = 7$  for plants, 16 for carabids and 21 for birds. In brackets are the number of times the considered landscape descriptor was selected by the random forest procedure. Detailed results on landscape descriptors selected for each biodiversity measure may be found in Appendices 1J-K. % PG: proportion of permanent grassland (%), PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges (km), % WH: proportion of wooded habitat (%), WH edges: wooded habitat edges (km), % crops: proportion of crop (%). “200m” and “500m” indicate the scale at which the considered landscape descriptor was measured.

Instead, the grassland assemblages of the three studied taxa were significantly influenced by wooded habitats. Random forest results showed that landscape descriptors related to wooded habitats were

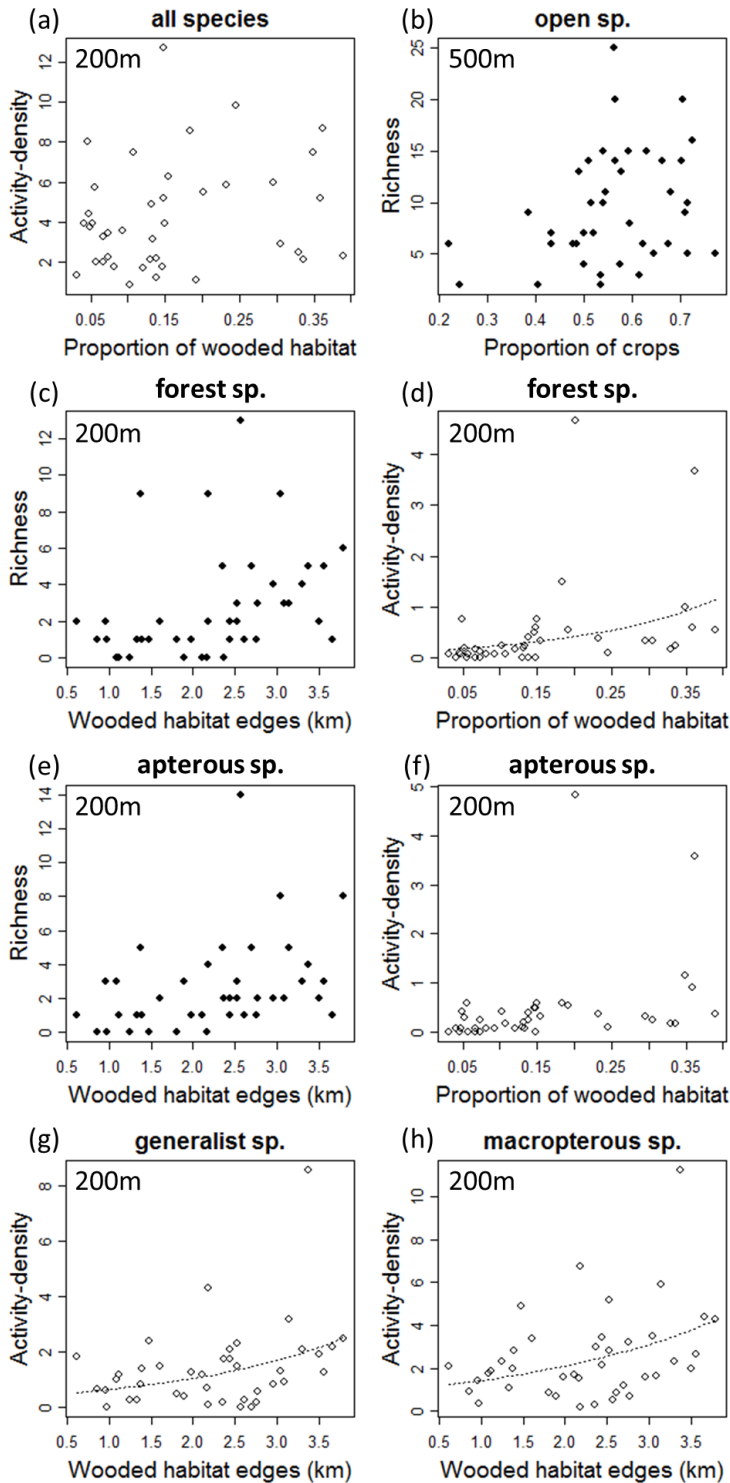
amongst those that explained the most variation in biodiversity measures, for the three taxa (Fig. 2). Significant relationships between landscape descriptors and biodiversity measures are shown graphically in Figures 3-5 and relationships that fell just short of the 95% confidence level are also displayed, but without a trend line. The latter relationships are henceforth referred to as nearly significant (near. sig.). Wooded habitats had a particularly strong influence at the 200m scale for birds and, to a lesser extent, carabid assemblages. Total plant species richness and richness of grassland plant specialists increased with increasing length of wooded habitat edges at the 200m scale (Fig. 3a/b, Appendix 1M). These two biodiversity measures responded similarly as most plant species were grassland specialists (Appendix 1B). The length of wooded habitat edges significantly positively influenced animal-dispersed plant species richness at the 200 m scale and gravity-dispersed plant species richness (near.sig.), at the 500 m scale (Fig. 3c/d, Appendix 1M). Ruderal and wind-dispersed plant species were unaffected by landscape descriptors.



**Figure 3.** Graphical representation of significant and nearly significant results for plants biodiversity measures. Barochorous: gravity-dispersed, zoochorous: animal-dispersed. Scale of effects is indicated on each graph corner as the radius of buffers surrounding sampled permanent grasslands. Plain lines are given for illustrative purposes (only for significant results). Coefficients from average models are used for drawing linear and exponential lines (Appendix 1M). Linear lines are used for biodiversity measures analysed with Gaussian distribution.

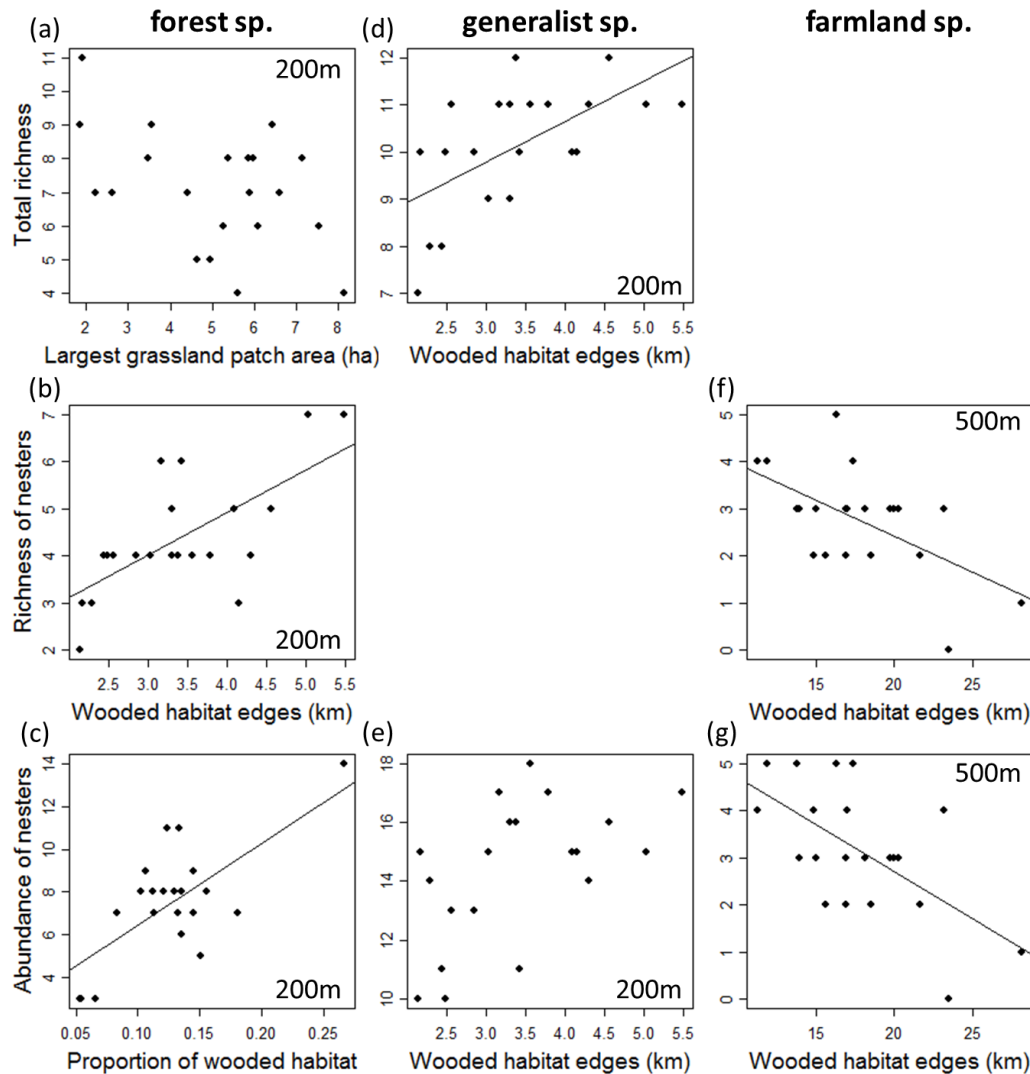
339 Total activity-density of carabid beetles tended to increase with increasing proportion of wooded  
340 habitats (near.sig. Fig. 4a, Appendix 1N). Richness of open habitat carabids increased with increasing  
341 proportion of crops area at the 500m scale (near. sig. Fig. 4b, Appendix 1N) which was strongly  
342 negatively correlated with the proportion of wooded habitats at this scale ( $r_s = -0.83$ , Appendix 1H).  
343 Species richness of forest carabid specialists increased with increasing length of wooded habitat edges  
344 and activity-density of the same group significantly increased with increasing proportion of wooded  
345 habitats, both at the 200m scale (Fig. 4c/d, Appendix 1N). Apterous carabid species, which are mostly  
346 forest specialists (Appendix 1C), followed the same trends (near. sig. Fig. 4e/f, Appendix 1N).  
347 Activity-density of carabid generalist and macropterous species were positively influenced by  
348 increasing length of wooded habitat edges, also at the 200m scale (Fig. 4g/h, Appendix 1N).





**Figure 4.** Graphical representation of significant and nearly significant results for carabid beetles biodiversity measures. Full dots are used for species richness, while empty dot and dashed lines are used for activity-density. Activity-density is expressed as number of individuals / valid trap / week. Scale of effects is indicated on each graph corner as the radius of buffers surrounding sampled permanent grasslands. Lines are given for illustrative purposes (only for significant results). Coefficients from average models are used for drawing linear and exponential lines (Appendix 1N). Exponential lines are used for Poisson and Negative Binomial distributions (log link functions).

358 For birds, richness of breeding forest specialists increased with increasing length of wooded habitat  
359 edges at the 200m scale while their abundance was positively influenced by proportion of wooded  
360 habitats also at the 200 m scale (Fig. 5b/c, Appendix 1O). As mentioned above, total richness of forest  
361 specialist bird species was the only biodiversity measure to be influenced by a grassland landscape  
362 descriptor, i.e a nearly significant negative influence of large grassland patch area (Fig 5a, Appendix  
363 1O). Similarly, total richness and abundance of breeding generalist birds increased with increasing  
364 length of wooded habitat edges at the 200m scale (Fig. 5d/e, Appendix 1O). Conversely, richness and  
365 abundance of breeding farmland bird specialists decreased with increasing length of wooded habitat  
366 edges, this time at the 500m scale (Fig. 5f/g, Appendix 1O). Birds with limited dispersal were not  
367 affected by landscape descriptors, while medium and high dispersal groups responded significantly  
368 (Appendix 1P) but this seems to have been more related to habitat affinity of the forest specialist or  
369 generalist species dominating these dispersal groups, than to dispersal ability per se, and so these  
370 results are not shown graphically.



**Figure 5.** Graphical representation of significant and nearly significant results for bird habitat affinity groups. Scale of effects is indicated on each graph corner as the radius of buffers surrounding sampled permanent grasslands. Lines are given for illustrative purposes (only for significant results). Coefficients from average models are used for drawing linear (Appendix 10), all models followed Gaussian distribution.

## 4. Discussion

The substantial variation in biodiversity between sampled permanent grasslands was not explained by the connectivity of permanent grassland, either amount or spatial configuration, in the landscape, at least at the two spatial scales considered (200 and 500 m). It should be kept in mind that overall diversity in such human-modified landscapes is low (see Irmiler & Hoernes, 2003), but our sampling protocol enabled us to detect a high proportion of the expected species pool for each taxonomic group. Many species were not considered to be grassland specialists, particularly in the animal groups, so the

lack of response of these assemblages as a whole is perhaps not surprising. However, contrary to our hypothesis, even grassland specialist groups were unaffected. The lack of response of grassland assemblages to grassland amount was also noted by Soderstrom et al. (2001), who found no effect on either species richness or composition of plants, ground beetles and other insects (butterflies, bumble bees, dung beetles), and birds . However, in their study the proportion of grassland varied only from 0 to 17%, while in this study it varied from at least 10 to 58% at the 200m scale and 4 to 32% at the 500m scale, and despite this, there was still little response from the different groups. In our study, grassland connectivity index (*sensu* Hanski 1999) also varied (0-7) without significantly influencing the grassland assemblages. Similarly, Öckinger et al. (2012) found grassland isolation (inverse of Hanski index) had no effect on either total species richness of plants and insects (butterflies, bees or hoverflies), or on richness of grassland specialist groups. One could argue that if we had used a wider range of connectivity indices we might have found different effects. However Villemey et al. (2015) explored a much wider range of connectivity measures including Hanski index but also nearest Euclidean distance, graph-based measures with cost (least-cost path) or resistance (circuit theoretic) distances, only to come to the same conclusion for the butterfly assemblages they studied. The only group to respond to any measure of grassland connectivity was the total species richness of forest birds, which included 6 species not breeding in our study area, while the richness of breeding forest species alone did not. This suggests that certain species with a greater affinity for forest habitats may avoid very open areas, even when engaged in more temporary activities than breeding, such as dispersal or foraging. Configuration of grassland at 500m was sometimes selected by random forest procedures for plant assemblages. However, as for the other taxa, the effects of these descriptors were not significant.

Grassland specialists dominated plant assemblages, forming on average 83% of total species richness. Both grassland and total species richness were positively influenced by wooded habitats in the landscape as also shown by Soderstrom et al. (2001) and Ernoult et al. (2006). One hypothesis is that landscapes with more wooded habitats tend to be less disturbed and more species rich, and that wooded edges, particularly hedgerows, have well-developed herbaceous strata which may act as

sources. Plant dispersal groups varied in their response to wooded habitats, independently from habitat affinity groups. Wind-dispersed species were unaffected by landscape context, in agreement with Piessens et al. (2005). However, we might have expected that wooded edges would inhibit flow of wind-dispersed seeds by forming physical barriers or reducing wind speed (Gaujour et al. 2012). Wooded habitat edges did promote animal-dispersed species, probably because seed-dispersing animals (birds, mammals, insects) tend themselves to follow woodland edges and hedgerows as they move.

In the case of carabids, the lack of grassland connectivity effects might be explained by the limited number of true grassland specialists (Roger et al. 2010; Neumann et al. 2016). Although grassland assemblages are clearly different from those observed in crop fields or wooded habitats (Duflot et al. 2015) some species considered to be crop or forest specialists are known to utilize permanent grasslands. Richness of open habitat carabid species tend to increase in landscapes with a greater proportion of crops, where a greater amount and diversity of complementary resources may be available (Duflot et al. 2016). Richness and activity-density of forest specialist species were higher in landscapes with more wooded habitat or edges (presence of hedgerows), which is a general observation for carabid assemblages found in farmland habitats (e.g. Millan-Pena et al. 2003; Aviron et al. 2005; Duflot et al. 2014). This is probably why total activity-density tended to increase with the proportion of wooded habitat in the landscape. Meanwhile, activity-density of generalist carabid species increased with increasing length of hedgerows, known to be overwintering sites for these species (Sotherton 1985; Thorbek and Bilde 2004). These results concur with studies in other contexts showing that carabid assemblages are strongly influenced by habitats adjacent to the focal habitat (Schneider et al. 2016; Yekwayo et al. 2016).

The majority of sampled birds were contacted in hedgerows delimiting each grassland or in trees or shrubs within, with relatively few observations of foraging in grasslands and < 1 breeding territory per hectare within the grassland habitat itself (data not shown). Therefore, it is not surprising that landscape descriptors of wooded habitat amount and configuration best explained distributions of

observed assemblages and more specifically of generalist and forest birds. Species more typical of open farmland habitats including grasslands are known to be negatively impacted by wooded or shrubby habitats in the landscape (Besnard et al. 2016) and we too found that wooded habitat edges reduced farmland bird richness and abundance of breeding farmland birds. Hedgerow avoidance behaviour of ground-nesting farmland birds is common and is partly due to increased predation risks associated with hedgerows and woodland edges (Besnard et al. 2016). The low dispersal ability group, which was composed of species with a diversity of habitat affinities, was not significantly affected by landscape context, although this group was expected to be the most sensitive to habitat connectivity. This study focused on breeding birds and we cannot rule out effects of landscape structure on this low dispersal group outside the breeding season and at wider geographical scales.

Given the composition of the sampled animal assemblages, it proved difficult to construct dispersal ability groups that would be independent from species habitat affinity and that would be comparable with other taxa. Although effects on plant dispersal groups were clearer, the positive, nearly significant, influence of wooded habitats on plant species dispersed by barochory was also difficult to explain. As Piessens et al. (2005) has highlighted, some plant species may disperse in a variety of ways making it difficult to define dispersal groups and to link dispersal processes with landscape structure.

Contrary to expectations, all three taxa responded mainly to landscape variables at the same 200 m scale. As the scale of landscape perception is expected to be between 4 to 9 times the average dispersal distance (Jackson and Fahrig 2012), the response of plant and carabids at the finer scale is less surprising than that of birds. It should be noted that most of the significant bird-landscape relationships related to breeding birds only, which are sedentary during the sampling season. Further investigation of a wider range of spatial scales of influence may yield different results. For both animal taxa, landscape configuration had opposite effects on open habitat and farmland species compared to forest and generalist species, and at different spatial scales. Farmland or open specialists seemed to have a wider scale of perception (500 m) than forest specialists and generalists (200 m), which suggests that

response to landscape context was more dependent on habitat affinity than taxonomic group. However, in the case of plants, ruderal species, more adapted to human disturbance and that included crop weeds, did not respond to increasing proportion of crops and/or decreased wooded habitats at the 500m scale. Interestingly, for both animal groups richness of forest species of tended to increase with connectivity of wooded habitats (i.e. hedgerows), while their abundance or activity-density increased with resource availability (i.e. amount of wooded habitat).

## 5. Conclusion

We found no evidence that increasing connectivity of common, mesophilic, permanent grassland would have positive or indeed negative effects on plant and animal assemblages of such grassland habitats. Hence conservation planning to enhance the surface area and linkages between such grasslands within agricultural landscapes alone is unlikely to produce biodiversity increases, at least as far as our three study taxa are concerned. Instead, studied assemblages responded mostly to wooded habitats surrounding sampled grasslands, including hedgerows. Lengths of wooded habitat edges were often more important than the amount of wooded habitat itself, although collinearity between these landscape descriptors made it difficult to disentangle their independent effects. For grassland plant assemblages as a whole, or for the generalist or forest specialist components of the grassland animal assemblages, more woodland habitat in the landscape matrix is generally positive. Preserving or re-creating landscapes composed of permanent grasslands interspersed with woodlands and hedgerows may be good policy, especially in regions where such forms of landscape organisation have been historically present and match with established and adapted biodiversity. Therefore, schemes aiming to reintroduce landscape complexity to farmland areas through hedge and tree planting should enhance biodiversity. However, the presence of open habitat carabids as well as farmland specialist birds depended on sufficiently large expanses of open land, free of wooded habitats. So, though increasing the area of common, permanent, mesophilic grassland did not influence these groups, maintaining sufficient areas of open land is crucial and potentially contradictory with the objectives of policies aiming to restore semi-natural, wooded habitats in farmland. These results illustrate the importance of

496 balancing landscape conservation planning in agricultural contexts, and of using multi-taxon  
497 approaches, to meet the needs of contrasting ecological groups.

498

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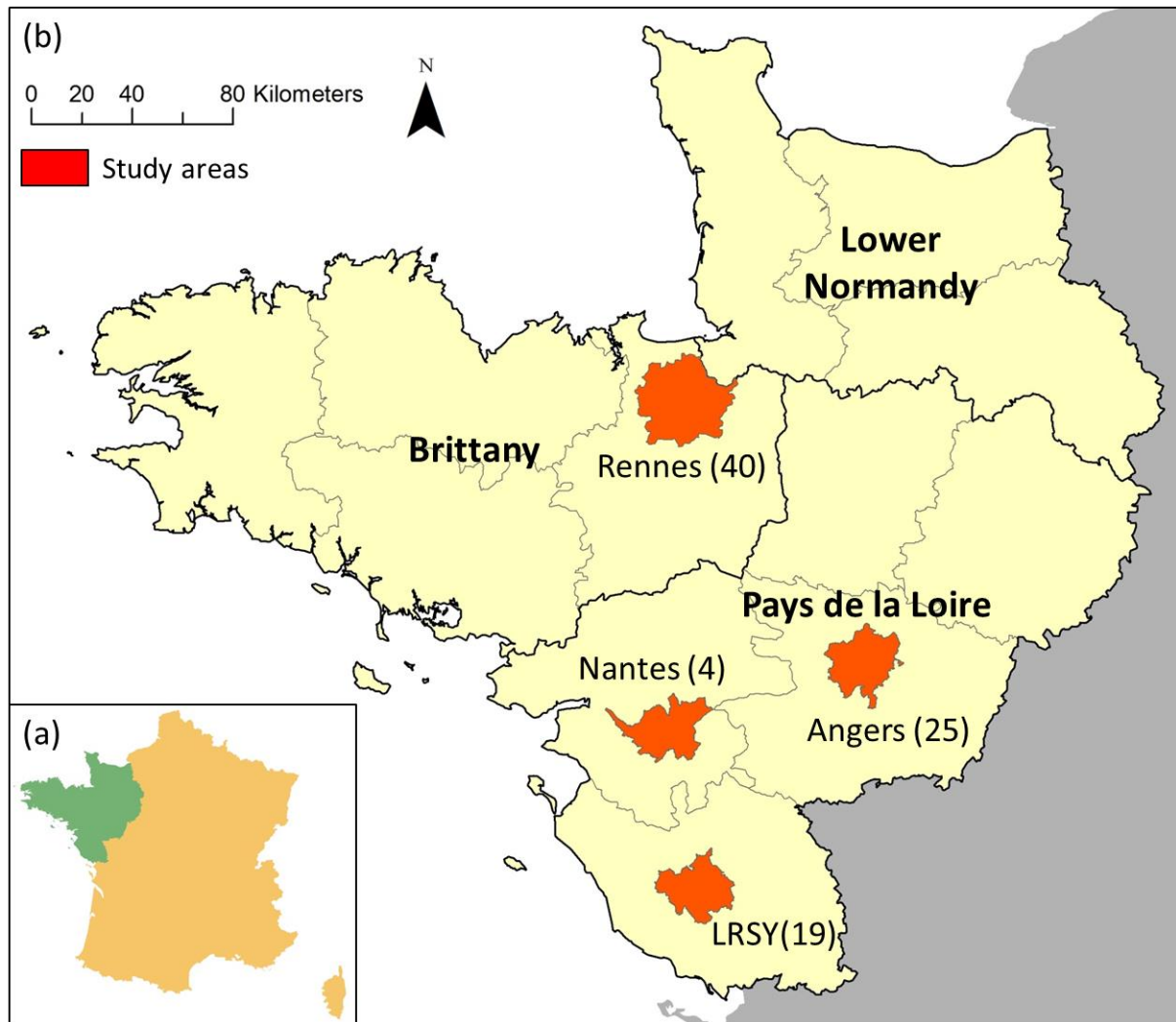
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700

## Online appendixes 1

### Appendix 1A

Map of the study areas. (a) Geographical location of the study region in France. (b) Geographical location of the four study areas in north-western France. In brackets, the number of sampled grasslands in each study area (LSRY = La Roche-sur-Yon).



## Appendix 1B

Distribution of plant species among the different biological groups within the sampled permanent grasslands. Species occurrence is the number of species observations across all sampled permanent grasslands (maximum occurrence = 55). Anemochorous: wind-dispersed, barochorous gravity-dispersed, zoochorous: animal-dispersed. NA: species for which data were not available, others: autochorous and hydrochorous.

Species number					
	anemochorous	barochorous	zoochorous	NA / others	<b>Total</b>
forest	3	4	6	1	<b>14</b>
grassland	10	25	23	1	<b>59</b>
ruderal	5	8	12	1	<b>26</b>
NA		1	2	6	<b>9</b>
<b>Total</b>	<b>18</b>	<b>38</b>	<b>43</b>	<b>9</b>	<b>108</b>

Summed species occurrences					
	anemochorous	barochorous	zoochorous	NA / others	<b>Total</b>
forest	6	11	21	1	<b>39</b>
grassland	131	243	296	2	<b>672</b>
ruderal	17	49	37	1	<b>104</b>
NA		1	2	7	<b>10</b>
<b>Total</b>	<b>154</b>	<b>304</b>	<b>356</b>	<b>11</b>	<b>825</b>



## Appendix 1C

Distribution of carabid beetle species among the different biological groups within the sampled permanent grasslands. Species occurrence is the number of species observations across all sampled permanent grasslands (maximum occurrence = 40). Apterous: windless, dimorphic: undeveloped wings, macropterous: winged. Activity-density is expressed as number of individuals / valid trap / week. NA: species for which data were not available.

Species number					
	apterous	dimorphic	macropterous	NA	Total
crop	2	2	11		<b>15</b>
forest	8	6	1		<b>15</b>
generalist	1	1	5		<b>7</b>
open habitat		7	11		<b>18</b>
NA	3	3	14	1	<b>21</b>
<b>Total</b>	<b>14</b>	<b>19</b>	<b>42</b>	<b>1</b>	<b>76</b>

Summed species occurrences					
	apterous	dimorphic	macropterous	NA	Total
crop	7	15	92		<b>114</b>
forest	79	29	3		<b>111</b>
generalist	14	1	152		<b>167</b>
open		184	194		<b>378</b>
NA	6	7	34	3	<b>50</b>
<b>Total</b>	<b>106</b>	<b>236</b>	<b>475</b>	<b>3</b>	<b>820</b>

Summed activity-density					
	apterous	dimorphic	macropterous	NA	Total
crop	0.70	1.62	15.20		<b>17.51</b>
forest	15.84	3.26	0.25		<b>19.35</b>
generalist	1.82	0.10	53.86		<b>55.77</b>
open habitat		44.96	27.90		<b>72.86</b>
NA	0.61	0.63	3.65	0.27	<b>5.16</b>
<b>Total</b>	<b>18.96</b>	<b>50.57</b>	<b>100.86</b>	<b>0.27</b>	<b>170.66</b>

## Appendix 1D

Distribution of bird species among the different biological groups within the sampled permanent grasslands, for total species pool and nesters. Species occurrence is the number of species observations across all sampled permanent grasslands (maximum occurrence = 21). Dispersal ability groups were obtained by splitting a dispersal ability predictor into categories using natural breaks (Jenks). NA: species for which data were not available. \* unstudied biological groups.

Species number						
	low	medium	high	long distance*	NA	Total
farmland	2	4	6	2	3	<b>17</b>
forest	5	8	6			<b>19</b>
generalist	6	4	6	2	2	<b>20</b>
urban*	3	1	5	2		<b>11</b>
NA					5	<b>5</b>
<b>Total</b>	<b>16</b>	<b>17</b>	<b>23</b>	<b>6</b>	<b>10</b>	<b>72</b>

Summed species occurrences						
	low	medium	high	long distance*	NA	Total
farmland	23	40	21	23	9	<b>116</b>
forest	29	90	61			<b>180</b>
generalist	112	78	35	33	2	<b>260</b>
urban*	23	7	22	16		<b>68</b>
NA					7	<b>7</b>
<b>Total</b>	<b>187</b>	<b>215</b>	<b>139</b>	<b>72</b>	<b>18</b>	<b>631</b>

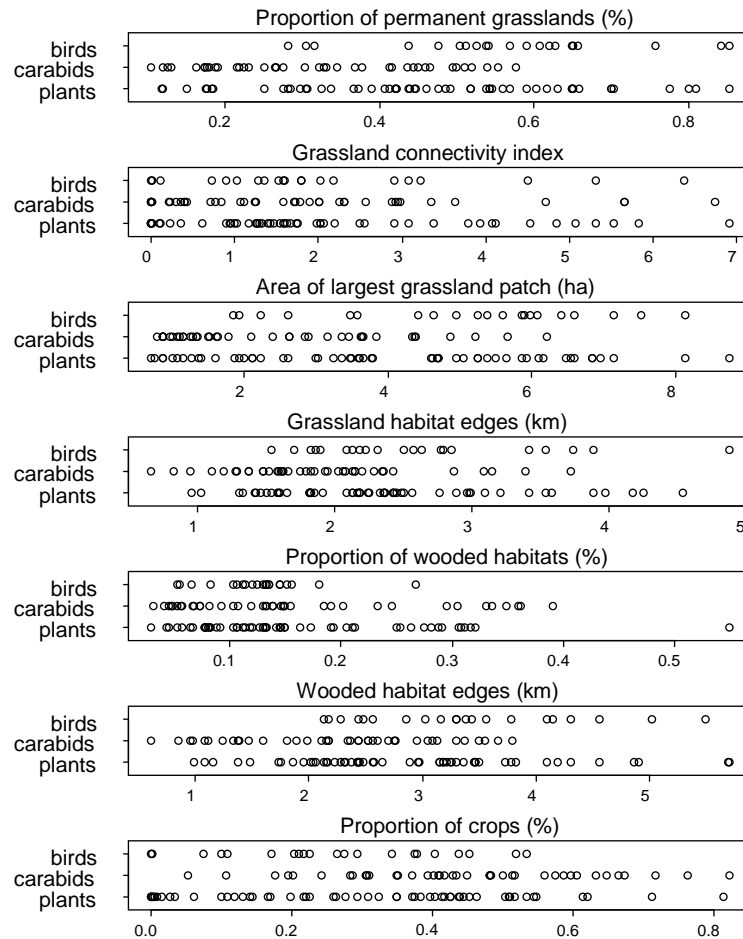
Nesting species number						
	low	medium	high	long distance*	NA	Total
farmland	2	4	3	2	1	<b>12</b>
forest	1	7	5			<b>13</b>
generalist	5	4	4	2		<b>15</b>
urban*	3	1	2			<b>6</b>
NA					1	<b>1</b>
<b>Total</b>	<b>11</b>	<b>16</b>	<b>14</b>	<b>4</b>	<b>2</b>	<b>47</b>

Summed nesting species occurrences						
	low	medium	high	long distance*	NA	Total
farmland	21	29	9	8	1	<b>68</b>
forest	24	65	25			<b>114</b>
generalist	98	66	25	18		<b>207</b>
urban*	13	4	7			<b>24</b>
NA					1	<b>1</b>
<b>Total</b>	<b>156</b>	<b>164</b>	<b>66</b>	<b>26</b>	<b>2</b>	<b>414</b>

Nesting species summed abundance						
	low	medium	high	long distance*	NA	Total
farmland	23	35	9	8	1	<b>76</b>
forest	42	126	30			<b>198</b>
generalist	182	131	37	21		<b>371</b>
urban*	15	4	7			<b>26</b>
NA					1	<b>1</b>
<b>Total</b>	<b>262</b>	<b>296</b>	<b>83</b>	<b>29</b>	<b>2</b>	<b>672</b>

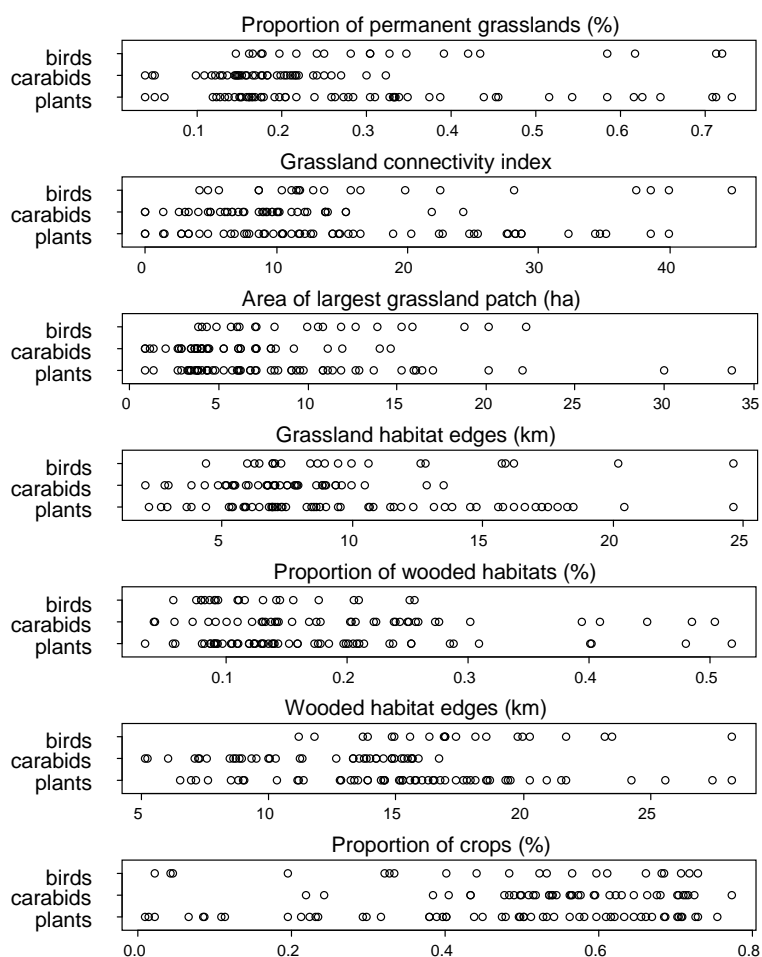
## Appendix 1E

Range of values of studied landscape descriptors, in a 200m-radius buffer surrounding sampled permanent grasslands, for each taxon. Each dot represents a landscape. Proportions are expressed as percentage.



## Appendix 1F

Range of values of studied landscape descriptors, in a 500m-radius buffer surrounding sampled permanent grasslands for each taxon. Each dot represents a landscape. Proportions are expressed as percentage.



## Appendix 1G

Pearson correlation coefficient for all pairs of landscape descriptors measured for plant-sampled permanent grasslands (n = 55). Highest correlation coefficients (> |0.7|) are indicated in bold characters. % PG: proportion of permanent grassland (%), PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges (km), % WH: proportion of wooded habitat (%), WH edges: wooded habitat edges (km), % crops: proportion of crop (%)

	200m							500m						
	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops	% PG	PG connect	PG LP	PG edges	% WH	WH edges	
200m														
% PG														
PG connect	0.39													
PG LP	0.79	-0.19												
PG edges	0.66	0.78	0.18											
% WH	-0.12	0.19	-0.20	0.19										
WH edges	0.13	0.38	-0.10	0.55	0.38									
% crops	-0.83	-0.49	-0.59	-0.70	-0.33	-0.29								
500m														
% PG	0.81	0.37	0.58	0.55	-0.02	0.21	-0.71							
PG connect	0.68	0.63	0.33	0.69	0.13	0.33	-0.69	0.87						
PG LP	0.70	0.05	0.68	0.21	-0.25	-0.03	-0.48	0.81	0.49					
PG edges	0.68	0.54	0.34	0.71	0.20	0.43	-0.74	0.88	0.88	0.53				
% WH	-0.18	0.09	-0.21	0.02	0.72	0.15	-0.22	-0.07	0.02	-0.26	0.13			
WH edges	0.36	0.33	0.18	0.48	0.34	0.74	-0.48	0.48	0.52	0.20	0.67	0.21		
% crops	-0.63	-0.37	-0.43	-0.52	-0.35	-0.31	0.79	-0.85	-0.78	-0.58	-0.88	-0.42	-0.57	

## Appendix 1H

Pearson correlation coefficient for all pairs of landscape descriptors measured for carabid-sampled permanent grasslands. (n = 40) Highest correlation coefficients (> |0.7|) are indicated in bold. % PG: proportion of permanent grassland (%), PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges (km), % WH: proportion of wooded habitat (%), WH edges: wooded habitat edges (km), % crops: proportion of crop (%)

	200m							500m						
	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops	% PG	PG connect	PG LP	PG edges	% WH	WH edges	
200m														
% PG														
PG connect	0.54													
PG LP	<b>0.76</b>	-0.09												
PG edges	0.64	<b>0.83</b>	0.14											
% WH	-0.03	0.33	-0.26	0.41										
WH edges	-0.10	0.28	-0.34	0.42	0.70									
% crops	<b>-0.80</b>	-0.69	-0.42	<b>-0.81</b>	-0.46	-0.23								
500m														
% PG	0.64	0.40	0.43	0.53	0.06	0.10	-0.51							
PG connect	0.42	<b>0.84</b>	-0.04	0.66	0.26	0.12	-0.54	0.48						
PG LP	0.52	-0.08	0.69	0.18	-0.16	-0.28	-0.27	0.66	-0.01					
PG edges	0.39	0.55	0.04	0.66	0.28	0.42	-0.45	<b>0.84</b>	0.63	0.29				
% WH	0.08	0.16	-0.03	0.12	0.49	0.28	-0.36	-0.08	0.02	-0.11	-0.09			
WH edges	-0.04	0.33	-0.30	0.41	0.60	<b>0.76</b>	-0.24	0.10	0.22	-0.29	0.46	0.28		
% crops	-0.42	-0.36	-0.21	-0.41	-0.42	-0.23	0.60	-0.45	-0.28	-0.31	-0.34	<b>-0.83</b>	-0.22	

## Appendix 1I

Pearson correlation coefficient for all pairs of landscape descriptors measured for bird-sampled permanent grasslands (n = 21). Highest correlation coefficients (> |0.7|) are indicated in bold characters. % PG: proportion of permanent grassland (%), PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges (km), % WH: proportion of wooded habitat (%), WH edges: wooded habitat edges (km), % crops: proportion of crop (%)

		200m							500m					
	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops	% PG	PG connect	PG LP	PG edges	% WH	WH edges	
200m														
% PG														
PG connect	0.49													
PG LP	0.60	-0.40												
PG edges	0.65	<b>0.73</b>	0.01											
% WH	0.23	0.46	-0.17	0.53										
WH edges	0.24	0.47	-0.19	0.63	<b>0.76</b>									
% crops	<b>-0.93</b>	-0.58	-0.45	<b>-0.75</b>	-0.42	-0.35								
500m														
% PG	<b>0.83</b>	0.53	0.38	0.62	0.18	0.24	<b>-0.84</b>							
PG connect	0.69	0.68	0.09	0.65	0.20	0.29	<b>-0.75</b>	<b>0.93</b>						
PG LP	<b>0.72</b>	0.08	0.69	0.35	0.05	0.01	-0.63	0.69	0.40					
PG edges	<b>0.76</b>	0.67	0.19	<b>0.76</b>	0.32	0.44	<b>-0.82</b>	<b>0.93</b>	<b>0.89</b>	0.55				
% WH	0.45	0.46	0.05	0.58	<b>0.72</b>	0.62	-0.60	0.58	0.55	0.29	0.70			
WH edges	0.28	0.52	-0.18	0.52	0.49	<b>0.70</b>	-0.39	0.38	0.42	0.04	0.64	<b>0.75</b>		
% crops	<b>-0.82</b>	-0.49	-0.42	-0.66	-0.30	-0.33	<b>0.88</b>	<b>-0.97</b>	<b>-0.88</b>	-0.67	<b>-0.92</b>	-0.69	-0.45	

## Appendix 1J

List of landscape descriptors retained for regression models after the random forest procedure for each plant biodiversity measure. The retained variable are those whose relative importance value was higher than the absolute importance value of the lowest negative-scoring landscape descriptor.

Biodiversity measure	200m							500m						
	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops
all species richness						X			X				X	
forest richness								X	X		X			X
grassland richness						X			X				X	
ruderal richness			X				X	X						
anemochorous richness														
barochorous richness											X		X	X
zoochorous richness						X					X			



## Appendix 1K

List of landscape descriptors retained for regression models after the random forest procedure for each carabid biodiversity measure. The retained variable are those whose relative importance value was higher than the absolute importance value of the lowest negative-scoring landscape descriptor.

Biodiversity measure	200m							500m						
	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops
all species richness														
all species activity-density					X									
crop richness														
forest richness				X	X	X								
generalist richness						X		X	X					
open richness												X		X
crop activity-density														
forest activity-density					X	X								X
generalist activity-density						X								
open activity-density														
apterous richness					X	X						X	X	X
dimorphic richness					X	X						X	X	X
macropterous richness			X									X		
apterous activity-density					X	X						X		X
dimorphic activity-density			X			X								
macropterous activity-density						X								

## Appendix 1L

List of landscape descriptors retained for regression models after the random forest procedure for each bird biodiversity measure. The retained variable are those whose relative importance value was higher than the absolute importance value of the lowest negative-scoring landscape descriptor.

Biodiversity measure	200m							500m						
	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops	% PG	PG connect	PG LP	PG edges	% WH	WH edges	% crops
all species total richness						X								
all species richness of nester					X	X								
all species abundance of nester					X	X						X		
forest total richness		X	X			X								X
generalist total richness					X	X						X	X	
farmland total richness								X	X					
forest richness of nester					X	X						X	X	
generalist richness of nester												X	X	
farmland richness of nester											X	X	X	
forest abundance of nester					X	X						X		
generalist abundance of nester					X	X						X	X	
farmland abundance of nester												X	X	
low richness														
medium richness			X		X									
high richness					X	X								
low richness of nester													X	
medium richness of nester														
high richness of nester				X	X	X								
low abundance of nester						X								
medium abundance of nester					X	X								
high abundance of nester					X	X								

## Appendix 1M

Estimate and 95% confidence intervals from model averaging for plant biodiversity measures. Significant results, i.e. estimates whose 95 % confidence interval do not include zero, are in bold. Near significant results are shown with a “X” sign. All models were performed using a Gaussian distribution except for biodiversity measures with \*, for which Poisson distribution was used, and with \*\* for which a negative binomial distribution was used. Models with only intercept values indicate a null model, i.e. random forest procedure selected no landscape descriptors (see “2.4 Statistical analyses”). Landscape descriptors: % PG: proportion of permanent grassland, PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges, % WH: proportion of wooded habitat, WH edges: wooded habitat edges, % crops: proportion of crop. “200m” and “500m” indicate the scale at which the considered landscape descriptor was measured.

biodiversity measure	landscape descriptor	estimates	lower CI	upper CI	
allspecies richness	(Intercept)	14,74	11,941	17,547	
	<b>WH edges 200m</b>	<b>2,05</b>	<b>0,123</b>	<b>3,983</b>	
	PG connect 500m	0,97	-0,734	2,678	
	WH edges 500m	0,64	-2,046	3,321	
forest richness *	(Intercept)	-0,42	-0,762	-0,074	
	PG edges 500m	0,33	-0,183	0,839	
	PG connect 500m	0,32	-0,180	0,821	
	% crops 500m	-0,25	-0,777	0,267	
	% PG 500m	0,17	-0,394	0,729	
grassland richness	(Intercept)	12,14	10,261	14,024	
	<b>WH edges 200m</b>	<b>1,97</b>	<b>0,488</b>	<b>3,443</b>	
	PG connect 500m	0,66	-0,667	1,986	
	WH edges 500m	0,12	-1,826	2,072	
ruderal richness **	(Intercept)	0,49	0,036	0,940	
	PG LP 200m	0,15	-0,091	0,400	
	% crops 200m	-0,15	-0,413	0,113	
	% PG 500m	0,02	-0,248	0,288	
anemochor richness *	(Intercept)	1,03	0,849	1,189	
barochor richness *	(Intercept)	1,67	1,451	1,887	
	WH edges 500m	0,15	-0,006	0,306	X
	% crops 500m	-0,08	-0,257	0,089	
	PG edges 500m	0,03	-0,195	0,248	
zoochor richness	(Intercept)	6,38	5,208	7,550	
	<b>WH edges 200m</b>	<b>1,36</b>	<b>0,577</b>	<b>2,134</b>	
	PG edges 500m	0,34	-0,461	1,142	

## Appendix 1N

Estimate and 95% confidence intervals from model averaging for carabid biodiversity measures. Significant results, i.e. estimates whose confidence interval do not include zero, are in bold. Near significant results are shown with a “X” sign. All models were performed using a negative binomial distribution except for biodiversity measures with \*, for which a Gaussian distribution was used and \*\* for which a Poisson distribution was used. Models with only intercept values indicate a null model, i.e. random forest procedure selected no landscape descriptors (see “2.4. Statistical analyses”). Landscape descriptors: % PG: proportion of permanent grassland, PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges, % WH: proportion of wooded habitat, WH edges: wooded habitat edges, % crops: proportion of crop. “200m” and “500m” indicate the scale at which the considered landscape descriptor was measured.

biodiversity measure	landscape descriptor	estimates	lower CI	upper CI	
allspecies richness	(Intercept)	3,02	2,890	3,153	
activity-density	(Intercept)	1,42	1,224	1,616	
	% WH 200m	0,18	-0,029	0,391	X
crop richness	(Intercept)	1,05	0,803	1,289	
activity-density **	(Intercept)	-0,98	-1,817	-0,514	
forest richness	(Intercept)	0,79	0,39	1,18	
	WH edges 200m	0,38	-0,02	0,78	X
	% WH 200m	0,33	-0,068	0,738	
	PG edges 200m	0,15	-0,235	0,534	
activity-density	(Intercept)	-1,08	-1,802	-0,367	
	<b>% WH 200m</b>	<b>0,56</b>	<b>0,015</b>	<b>1,108</b>	
	% crops 500m	-0,42	-1,050	0,203	
	WH edges 200m	0,45	-0,309	1,200	
generalist richness *	(Intercept)	4,18	3,456	4,894	
	WH edges 200m	0,45	-0,262	1,155	
	PG connect 500m	-0,36	-1,087	0,366	
	% PG 500m	-0,11	-0,839	0,629	
activity-density	(Intercept)	0,18	-0,210	0,561	
	<b>WH edges 200m</b>	<b>0,42</b>	<b>0,078</b>	<b>0,765</b>	
open richness	(Intercept)	2,23	2,050	2,410	
	% crops 500m	0,20	-0,028	0,435	X
	% WH 500m	-0,13	-0,416	0,157	
activity-density	(Intercept)	0,55	0,204	0,814	
apterous richness	(Intercept)	0,82	0,490	1,147	
	WH edges 200m	0,31	-0,064	0,692	X
	% WH 500m	0,24	-0,113	0,592	
	% WH 200m	0,27	-0,112	0,648	
	% crops 500m	-0,21	-0,563	0,147	
	WH edges 500m	0,23	-0,178	0,643	

activity-density	(Intercept)	-1,11	-1,834	-0,389	X
	% WH 200m	0,51	-0,063	1,089	
	% crops 500m	-0,48	-1,147	0,192	
	WH edges 200m	0,40	-0,343	1,151	
	% WH 500m	0,30	-0,513	1,112	
dimorphic richness *	(Intercept)	5,90	4,737	7,063	
	WH edges 200m	-1,38	-3,003	0,234	
	% crops 500m	0,94	-0,700	2,587	
	% WH 500m	-0,62	-2,479	1,243	
	WH edges 500m	-0,76	-2,568	1,058	
activity-density	% WH 200m	-0,16	-2,069	1,741	
	(Intercept)	0,11	-0,279	0,505	
	WH edges 200m	-0,31	-0,649	0,023	
	PG LP 200m	0,06	-0,295	0,405	
	(Intercept)	2,47	2,293	2,645	
macropterous richness	PG LP 200m	-0,14	-0,323	0,049	
	% WH 500m	-0,10	-0,290	0,081	
	(Intercept)	0,84	0,583	1,105	
activity-density	WH edges 200m	<b>0,34</b>	<b>0,086</b>	<b>0,586</b>	

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## Appendix 10

Estimate and 95% confidence intervals from model averaging for bird biodiversity measures (all species and habitat affinity). Significant results, i.e. estimates whose 95 % confidence interval do not include zero, are in bold. Near significant results are shown with a “X” sign. All models were performed using a Gaussian distribution except for biodiversity measures with \*, for which a Poisson distribution was used. Models with only intercept values indicate a null model, i.e. random forest procedure selected no landscape descriptors (see “2.4. Statistical analyses”). Landscape descriptors: % PG: proportion of permanent grassland, PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges, % WH: proportion of wooded habitat, WH edges: wooded habitat edges, % crops: proportion of crop. “200m” and “500m” indicate the scale at which the considered landscape descriptor was measured.

biodiversity measure	Landscape descriptor	estimates	lower CI	upper CI	
all species total richness	(Intercept)	24,29	22,623	25,949	
	WH edges 200m	1,18	-0,508	2,863	
richness of nester	(Intercept)	15,87	14,869	16,865	
	% WH 200m	0,59	-0,270	1,459	
abundance of nester	WH edges 200m	0,52	-0,381	1,431	
	(Intercept)	25,81	24,27	27,35	
	% WH 200m	2,01	-0,31	4,32	
	WH edges 200m	1,60	-0,76	3,95	
	% WH 500m	-0,39	-2,70	1,92	
forest total richness	(Intercept)	7,10	6,345	7,845	
	PG LP 200m	-0,76	-1,526	0,009	X
	WH edges 200m	0,66	-0,155	1,483	
	PG connect 200m	0,52	-0,338	1,381	
	WH edges 500m	0,30	-0,666	1,269	
richness of nester	(Intercept)	4,38	3,934	4,827	
	<b>WH edges 200m</b>	<b>0,85</b>	<b>0,238</b>	<b>1,456</b>	
	% WH 200m	0,74	-0,018	1,504	X
	% WH 500m	-0,30	-0,947	0,349	
	WH edges 500m	-0,19	-0,840	0,455	
abundance of nester	(Intercept)	7,43	5,151	9,710	
	<b>% WH 200m</b>	<b>1,76</b>	<b>0,786</b>	<b>2,734</b>	
	WH edges 200m	0,38	-0,79	1,56	
	% WH 500m	-0,25	-1,432	0,924	
generalist total richness	(Intercept)	10,14	9,639	10,647	
	<b>WH edges 200m</b>	<b>0,81</b>	<b>0,111</b>	<b>1,501</b>	
	% WH 200m	0,62	-0,099	1,330	
	WH edges 500m	-0,44	-1,148	0,268	
	% WH 500m	-0,03	-0,699	0,647	
richness of nester*	(Intercept)	2,08	1,924	2,246	
	WH edges 500m	0,05	-0,116	0,210	
	% WH 500m	0,04	-0,118	0,207	

abundance of nester	(Intercept)	14,48	13,50	15,46	X
	WH edges 200m	1,19	-0,02	2,40	
	% WH 200m	0,63	-0,90	2,15	
	% WH 500m	0,67	-0,68	2,01	
	WH edges 500m	0,27	-1,269	1,818	
<hr/>					
farmland total richness*	(Intercept)	1,55	1,338	1,760	
	PG connect 500m	0,09	-0,116	0,300	
	% PG 500m	0,08	-0,132	0,287	
richness of nester	(Intercept)	2,76	2,330	3,194	
	<b>WH edges 500m</b>	<b>-0,62</b>	<b>-1,096</b>	<b>-0,153</b>	
	% WH 500m	-0,40	-1,058	0,250	
	PG edges 500m	-	-	-	
abundance of nester	(Intercept)	3,14	2,631	3,655	
	<b>WH edges 500m</b>	<b>-0,82</b>	<b>-1,389</b>	<b>-0,248</b>	
	% WH 500m	-0,40	-1,282	0,478	

## Appendix 1P

Estimate and 95% confidence intervals from model averaging for bird biodiversity measures (dispersal ability). Significant results, i.e. estimates whose confidence interval do not include zero, are in bold. All models were performed using a Gaussian distribution. Models with only intercept values indicate a null model, i.e. random forest procedure selected no landscape descriptors (see “2.4. Statistical analyses”). Landscape descriptors: % PG: proportion of permanent grassland, PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges, % WH: proportion of wooded habitat, WH edges: wooded habitat edges, % crops: proportion of crop. “200m” and “500m” indicate the scale at which the considered landscape descriptor was measured.

Biodiversity measure	landscape descriptor	estimates	lower CI	upper CI
low total richness	(Intercept)	6.81	6.281	7.338
richness of nester	(Intercept)	5.76	5.297	6.230
	WH edges 500m	0.36	-0.058	0.781
abundance of nester	(Intercept)	9.81	8.970	10.649
	WH edges 200m	0.66	-0.184	1.501
medium total richness	(Intercept)	8.43	7.754	9.104
	PG LP 200m	-0.66	-1.332	0.013
	% WH 200m	0.39	-0.314	1.098
richness of nester	(Intercept)	6.33	5.828	6.906
abundance of nester	(Intercept)	11.43	10.339	12.518
	% WH 200m	<b>1.38</b>	<b>0.087</b>	<b>2.682</b>
	WH edges 200m	0.08	-1.811	1.974
high total richness	(Intercept)	5.48	4.820	6.132
	WH edges 200m	0.52	-0.276	1.323
	% WH 200m	-0.07	-1.078	0.942
richness of nester	(Intercept)	2.62	2.129	3.109
	WH edges 200m	<b>0.67</b>	<b>0.046</b>	<b>1.292</b>
	% WH 200m	0.62	-0.004	1.254
	PG edges 200m	-0.19	-0.838	0.467
abundance of nester	(Intercept)	3.33	2.588	4.078
	% WH 200m	0.85	-0.060	1.761
	WH edges 200m	0.76	-0.217	1.745



## Appendix 2A

Corrected Akaike information criterion ( $AIC_c$ ),  $AIC_c$  difference with best model ( $\Delta AIC_c$ ), and Akaike weights ( $w_i$ ) of all supported models ( $\Delta AIC_c < 4$ ) that were included in model averaging for the analysis of plant biodiversity measures. Models are ordered in terms of  $\Delta AIC_c$ . In bold are the biodiversity measures with significant effects detected. % PG: proportion of permanent grassland (%), PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges (km), % WH: proportion of wooded habitat (%), WH edges: wooded habitat edges (km), % crops: proportion of crop (%)

Biodiversity measure	Intercept	PG LP 200m	WH edges 200m	% crops 200m	% PG 500m	PG connect 500m	PG edges 500m	WH edges 500m	% crops 500m	$AIC_c$	$\Delta AIC_c$	$w_i$
<b>allspecies richness</b>	14.77		2.03			0.99				345.7	0	0.25
	14.76		1.99			0.97		0.08		345.9	0.20	0.23
	14.69		1.92					0.61		346.1	0.45	0.20
	14.76		2.30							346.2	0.47	0.20
	14.75					0.90		1.56		348.6	2.88	0.06
	14.68							2.02		348.7	3.04	0.06
forest richness	-0.42						0.38			119.6	0	0.19
	-0.42					0.38				119.7	0.04	0.19
	-0.41								-0.37	120.3	0.68	0.14
	-0.40				0.33					120.9	1.25	0.10
	-0.43					0.25			-0.17	121.6	1.93	0.07
	-0.43					0.20	0.21			121.6	1.94	0.07
	-0.42						0.28		-0.11	121.9	2.21	0.06
	-0.42				0.04		0.34			122.0	2.31	0.06
	-0.42				0.05	0.34				122.0	2.35	0.06
	-0.42				0.12				-0.26	122.4	2.80	0.05

<b>grassland richness</b>	12.15		2.08				320.5	0	0.33
	12.15		1.89		0.65		321.0	0.46	0.26
	12.12		1.90			0.28	321.2	0.66	0.23
	12.15		1.94		0.67	-0.08	321.7	1.17	0.18
ruderal richness	0.50						190.7	0	0.24
	0.49	0.17					190.8	0.13	0.23
	0.47		-0.16				191.0	0.32	0.21
	0.50			0.08			192.4	1.75	0.10
	0.48	0.11	-0.10				192.7	2.06	0.09
	0.49	0.17		0.00			193.2	2.55	0.07
	0.47		-0.20	-0.05			193.3	2.63	0.07
anemochor richness	1.03						199.7		
<b>barochor richness</b>	1.66				0.16		249.6	0	0.38
	1.66				0.13	-0.05	251.5	1.88	0.15
	1.68					-0.11	251.7	2.08	0.13
	1.66			0.02	0.15		251.9	2.27	0.12
	1.69			0.10			252.4	2.81	0.09
	1.70						252.7	3.11	0.08
	1.66			-0.08	0.15	-0.11	253.6	3.94	0.05
<b>zoochor richness</b>	6.38		1.39				268.5	0	0.70
	6.38		1.26		0.34		270.2	1.71	0.30

## Appendix 2B

Corrected Akaike information criterion ( $AIC_c$ ),  $AIC_c$  difference with best model ( $\Delta AIC_c$ ), and Akaike weights ( $w_i$ ) of all supported models ( $\Delta AIC_c < 4$ ) that were included in model averaging for the analysis of carabid biodiversity measures. Models are ordered in terms of  $\Delta AIC_c$ . In bold are the biodiversity measures with significant effects detected. % PG: proportion of permanent grassland (%), PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges (km), % WH: proportion of wooded habitat (%), WH edges: wooded habitat edges (km), % crops: proportion of crop (%)

Biodiversity measure	Intercept	PG LP 200m	PG edges 200m	% WH 200m	WH edges 200m	% PG 500m	PG connect 500m	% WH 500m	WH edges 500m	% crops 500m	$AIC_c$	$\Delta AIC_c$	$w_i$
allspecies richness	3.02										288.7		
<b>activity-density</b>	1.41			0.18							194.2	0	0.65
	1.43										195.4	1.26	0.35
crop richness	1.05										173.5		
activity-density	-0.98										69.3		
<b>forest richness</b>	0.78				0.43						174.3	0	0.33
	0.80			0.40							174.9	0.61	0.24
	0.78			0.22	0.28						176.0	1.68	0.14
	0.78		0.11		0.38						176.6	2.28	0.10
	0.80		0.14	0.35							177.0	2.65	0.09
	0.79										178.0	3.71	0.05
	0.79		0.27								178.2	3.94	0.05
<b>activity-density</b>	-1.06			0.61							76.5	0	0.33
	-1.09			0.49						-0.30	78.1	1.67	0.14
	-1.10				0.60						78.5	2.00	0.12
	-1.11									-0.55	78.5	2.01	0.12

	-1.16		0.49			-0.44	78.6	2.12	0.12
	-1.09	0.50	0.22				78.8	2.30	0.11
	-1.00						80.0	3.57	0.06
generalist richness	4.18						179.5	0	0.38
	4.18		0.43				180.7	1.22	0.21
	4.18			-0.34			181.3	1.82	0.15
	4.18			-0.09			182.1	2.63	0.10
	4.18		0.48	-0.40			182.3	2.81	0.09
	4.18		0.45	-0.14			183.4	3.91	0.05
activity-density	0.17		0.42				124.7	0	1.00
open richness	2.23					0.21	246.5	0	0.47
	2.23				-0.18		247.7	1.26	0.25
	2.24						248.7	2.25	0.15
	2.23				-0.03	0.18	249.0	2.59	0.13
activity-density	0.55						142.7		
apterous richness	0.81		0.33		0.24		168.8	0	0.10
	0.85	0.38					169.0	0.21	0.09
	0.82		0.40				169.1	0.25	0.09
	0.81		0.35			-0.23	169.1	0.27	0.09
	0.80				0.27	0.29	169.7	0.91	0.06
	0.83	0.23	0.24				170.2	1.37	0.05
	0.83	0.29			0.19		170.2	1.39	0.05
	0.81					0.31	170.2	1.39	0.05
	0.80				0.35		170.3	1.46	0.05
	0.84	0.31				-0.17	170.4	1.56	0.04
	0.84	0.29				0.19	170.5	1.71	0.04
	0.82					0.37	170.8	1.94	0.04
	0.81					-0.32	171.3	2.46	0.03
	0.80		0.24		0.24	0.12	171.3	2.50	0.03
	0.81	0.11	0.26		0.20		171.3	2.50	0.03

activity-density	0.82		0.31		0.13		171.3	2.54	0.03
	0.82	0.14	0.26			-0.19	171.4	2.61	0.03
	0.81		0.33	0.16		-0.09	171.5	2.64	0.03
	0.81		0.26		0.12	-0.23	171.6	2.74	0.02
	0.82	0.18		0.20	0.20		171.7	2.90	0.02
	0.82	0.20			0.20	-0.18	171.8	3.04	0.02
	0.80			0.20	0.30	-0.09	172.4	3.58	0.02
	0.83	0.22	0.18		0.10		172.8	3.95	0.01
	-1.07	0.60					75.6	0	0.20
	-1.13					-0.59	76.6	0.98	0.12
	-1.12	0.46				-0.37	76.6	0.98	0.12
	-1.17		0.46			-0.50	77.1	1.42	0.10
	-1.11	0.49		0.25			77.6	1.94	0.07
	-1.13			0.54			77.6	1.94	0.07
	-1.10		0.57				77.7	2.10	0.07
	-1.09	0.50	0.19				78.0	2.39	0.06
	-1.16		0.45	0.40			78.3	2.66	0.05
	-1.02						78.9	3.28	0.04
	-1.14	0.32	0.22			-0.39	79.1	3.45	0.04
	-1.14			0.09		-0.51	79.2	3.57	0.03
	-1.11	0.47		-0.10		-0.45	79.3	3.71	0.03
dimorphic richness	5.90		-1.45			1.00	220.9	0	0.10
	5.90		-1.42	-0.91			221.5	0.56	0.07
	5.90		-1.42	-0.22		0.81	221.7	0.81	0.06
	5.90		-1.16		-0.38	0.98	221.9	1.02	0.06
	5.90		-1.68				222.0	1.04	0.06
	5.90	0.15	-1.54			1.04	222.1	1.22	0.05
	5.90				-1.25	1.05	222.5	1.55	0.04
	5.90		-1.15	-0.88	-0.36		222.5	1.58	0.04
	5.90	0.17	-1.53	-0.96			222.6	1.71	0.04

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## Appendix 2C

Corrected Akaike information criterion ( $AIC_c$ ),  $AIC_c$  difference with best model ( $\Delta AIC_c$ ), and Akaike weights ( $w_i$ ) of all supported models ( $\Delta AIC_c < 4$ ) that were included in model averaging for the analysis of bird biodiversity measures. Models are ordered in terms of  $\Delta AIC_c$ . In bold are the biodiversity measures with significant effects detected. % PG: proportion of permanent grassland (%), PG connect: permanent grassland connectivity index, PG LP: largest permanent grassland patch (ha), PG edges: permanent grassland edges (km), % WH: proportion of wooded habitat (%), WH edges: wooded habitat edges (km), % crops: proportion of crop (%)

Biodiversity measure	Intercept	PG LP 200m	PG edges 200m	% WH 200m	WH edges 200m	% PG 500m	PG connect 500m	% WH 500m	WH edges 500m	$AIC_c$	$\Delta AIC_c$	$w_i$
all species total richness	24.29				1.18					118.8	0	0.55
	24.29									119.3	0.44	0.45
richness of nester	15.86									88.0	0	0.35
	15.87			0.64						88.2	0.24	0.31
	15.87				0.60					88.5	0.59	0.26
	15.87			0.42	0.28					90.8	2.86	0.08
abundance of nester	25.81			2.23						115.7	0	0.28
	25.81			1.42	1.07					116.2	0.52	0.21
	25.81				2.15					116.4	0.68	0.20
	25.81			2.57				-0.47		117.0	1.34	0.14
	25.81			1.81	1.19			-0.66		117.8	2.15	0.09
	25.81				2.11			0.06		118.1	2.41	0.08
<b>forest total richness</b>	7.10		-0.82							88.2	0	0.24
	7.10		-0.71		0.60					89.0	0.82	0.16
	7.10				0.74					89.2	1.00	0.14
	7.10									89.8	1.57	0.11

	7.10	0.66				90.0	1.78	0.10
	7.10	0.40	-0.66			90.7	2.52	0.07
	7.10		-0.76		0.35	91.0	2.77	0.06
	7.10				0.49	91.4	3.15	0.05
	7.10	0.40		0.55		91.7	3.48	0.04
	7.10			0.78	-0.06	92.2	3.96	0.03
<b>richness of nester</b>	4.38			0.88		68.9	0	0.44
	4.38		0.84			70.3	1.40	0.22
	4.38		0.41	0.57		71.2	2.32	0.14
	4.38			1.01	-0.22	72.5	3.63	0.07
	4.38			1.01	-0.19	72.5	3.65	0.07
	4.38		1.12		-0.39	72.7	3.83	0.06
<b>abundance of nester</b>	7.43			1.78		90.4	0	0.64
	7.43			1.49	0.38	92.7	2.37	0.19
	7.43			1.98	-0.25	93.0	2.66	0.17
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<b>generalist total richness</b>	10.14			0.78		73.6	0	0.44
	10.14		0.73			74.7	1.12	0.25
	10.14			1.09	-0.44	75.7	2.13	0.15
	10.14		0.34	0.52		76.4	2.81	0.11
	10.14			0.80	-0.03	77.6	3.98	0.06
richness of nester	2.09					90.0	0	0.62
	2.08				0.05	92.4	2.38	0.19
	2.08				0.04	92.5	2.42	0.19
<b>abundance of nester</b>	14.48			1.25		97.8	0	0.33
	14.48		1.03			100.0	2.21	0.11
	14.48		0.21	1.09		100.0	2.26	0.11
	14.48			1.01	0.39	100.1	2.29	0.11
	14.48				1.01	100.2	2.40	0.10
	14.48			1.29	-0.05	100.3	2.51	0.09
	14.48				0.85	101.4	3.63	0.05



	14.48				101.5	3.76	0.05
	14.48	0.64		0.56	101.7	3.95	0.05
farmland total richness	1.55				85.5	0	0.58
	1.55		0.09		87.4	1.90	0.22
	1.55	0.08			87.7	2.15	0.20
richness of nester	2.76			-0.64	66.8	0	0.64
	2.76			-0.54	69.3	2.56	0.18
	2.76			-0.15 -0.53	70.6	3.78	0.10
	2.76				70.7	3.92	0.09
abundance of nester	3.14			-0.83	73.9	0	0.74
	3.14			-0.13 -0.74	77.4	3.52	0.13
	3.14			-0.68	77.4	3.53	0.13