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Arthropod traits and assemblages differ between core patches, transient stepping-stones and landscape corridors.

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Abstract

Context

Restoring landscape connectivity can mitigate fragmentation and improve population resilience, but functional equivalence of contrasting elements is poorly understood. Evaluating biodiversity outcomes requires examining assemblage-responses across contrasting taxa.

Objectives

We compared arthropod species and trait composition between contrasting open-habitat network elements: core patches, corridors (allowing individual dispersal and population percolation), and transient stepping-stones (potentially enhancing meta-population dynamics).

Methods

Carabids and spiders were sampled from core patches of grass-heath habitat (n=24 locations across eight sites), corridors (trackways, n=15) and recently-replanted clear-fells (transient patches, n=19) set in a forest matrix impermeable to open-habitat arthropods. Species and trait (habitat association, diet, body size, dispersal ability) composition were compared by ordination and fourth corner analyses.

Results

Each network element supported distinct arthropod assemblages with differing functional trait composition. Core patches were dominated by specialist dry-open habitat species while generalist and woodland species contributed to assemblages in connectivity elements. Nevertheless, transient patches (and to a lesser degree, corridors) supported dry-open species characteristic of the focal grass-heath sites. Trait associations differed markedly among the three elements. Dispersal mechanisms and their correlates differed between taxa, but dry-open species in transient patches were characterised by traits favouring dispersal (large running hunter spiders and large, winged, herbivorous carabids), in contrast to wingless carabids in corridors.

Conclusions

Core patches, dispersal corridors and transient stepping-stones are not functionally interchangeable within this system. Semi-natural core patches supported a filtered subset of the regional fauna. Evidence for enhanced connectivity through percolation (corridors) or meta-population dynamics (stepping stones) differed between the two taxa.

Keywords

Dispersal corridors, Ecological network, Landscape connectivity, Movement corridors, Open-habitat network

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1 Introduction

2 Restoring functional connectivity is promoted in conservation strategies to facilitate
3 biodiversity resilience and population survival in the face of anthropogenic landscape
4 fragmentation, land-use intensification and climate change (Chetkiewicz et al. 2006, DEFRA
5 2018, Isaac et al. 2018). At local scales, enhanced connectivity within regional landscapes
6 may improve local population persistence by enhancing rescue effects and reducing
7 detrimental impacts of fragmentation (Beier and Noss 1998, Damschen et al. 2006) while at
8 greater scales, connectivity is advocated to facilitate population range-response to climatic
9 change (e.g. Heller and Zavaleta 2009, Imbach et al. 2013). While the optimal level of site
10 aggregation or dispersion may differ between the aims of enhancing local persistence or
11 favouring large-scale range-expansion (Hodgson et al. 2011), functional connectivity at both
12 scales requires appropriate interventions within local landscapes (Jongman et al. 2004,
13 Jongman et al. 2011). Within such local networks, although some generalist species may
14 disperse through surrounding matrix habitats, for specialist species population connectivity
15 often requires connectivity elements of suitable quality and type, with corridors, stepping
16 stones, or a less hostile matrix (Bennett 2003) intended to improve dispersal and functional
17 connectivity between 'core patches' or 'ecological refuges' — relict fragments of natural or
18 semi-natural habitat that have retained local species-populations over decadal timescales
19 (Davis et al. 2013). Despite the abundance of literature promoting restoration of ecological
20 networks the relative benefits of corridor linkages or stepping stones are poorly understood
21 (Dolman 2012). Furthermore, few connectivity studies report community or assemblage
22 responses, with research often focused on a few target species (Haddad et al. 2014).
23 Conservation strategies seeking to restore ecological connectivity are best informed by
24 understanding responses to connectivity elements of differing configurations across

25 multiple taxa (Pedley et al. 2013b, Kormann et al. 2015, Albert et al. 2017). Additionally,
26 examining responses in terms of assemblages' trait- rather than species-composition
27 supports generalisation to other ecological systems (McGill et al. 2006, Pedley and Dolman
28 2014, Santini et al. 2016).

29 Connectivity is often conceived as movement corridors enabling dispersal between primary
30 habitat patches (Simberloff et al. 1992). Meta-analysis of movement, considering both direct
31 (movement frequency and rate) and indirect (abundance and species richness in connected
32 patches) measures, confirms corridors increase species' movement in fragmented
33 landscapes (Gilbert-Norton et al. 2010), potentially recolonising vacant habitat or enhancing
34 persistence through population rescue (Lawson et al. 2012). For slowly-dispersing and
35 matrix-sensitive species, corridors of suitable quality may allow population percolation over
36 multiple generations (Bennett 2003), assisting range responses to climate change (Krosby et
37 al. 2010). However, corridors, which by their nature are narrow with high edge-area ratios
38 (Simberloff et al. 1992), may have assemblages filtered and modified by edge impacts
39 (Ewers and Didham 2008, Campbell et al. 2011). Alternatively, stepping stones, a series of
40 discrete, smaller patches intended to link otherwise isolated habitat blocks, may provide
41 connectivity for mobile species capable of occasional dispersal across the intervening matrix
42 (Schultz 1998). A reduced edge-area may mitigate edge impacts and offer better habitat
43 suitability for specialist species, although assemblages may be filtered by colonisation ability
44 (Kormann et al. 2015). Both of these types of connectivity elements are often advocated
45 without explicit consideration of their relative efficacy or functional equivalence (e.g.
46 Lawton et al. 2010), with limited analysis of the trade-offs between differing configurations
47 and taxa.

48 For species associated with early-successional or physically-disturbed habitats, stepping-
49 stone suitability may be short-lived, with more frequent dispersal required for meta-
50 population persistence (Amarasekare and Possingham 2001, Johst et al. 2002, Loehle 2007).
51 Similarly, cyclic or episodic landuse, including much habitat management practiced for
52 conservation, the development and building phases of physical infrastructure projects, and
53 crop or forestry rotations, may all provide short-lived opportunities for specialists able to
54 colonise, exploit and move on from stepping stones. Understanding assemblage responses
55 to differing configuration of habitat can inform whether biodiversity conservation strategies
56 can take advantage of spatially discontinuous and transient patches, or instead require
57 permanent and spatially continuous, connectivity elements.

58 Here, we examine arthropod (carabid and spider) trait and assemblage responses to open-
59 habitat elements of contrasting configuration, comprising: 'core patches' (semi-natural
60 grass-heath sites supporting large numbers of scarce and threatened taxa), grass-heath
61 habitat persisting as 'linear corridors' (formed by trackways and verges), and transient
62 stepping-stone patches (formed by infrequent systematic physical disturbance events), with
63 the latter two potential connectivity elements embedded in a plantation matrix
64 impermeable to open habitat specialists (Pedley 2012). A fundamental assumption of our
65 study, is that assemblage richness and composition in the recently-created connecting
66 elements has been shaped by patterns of individual dispersal and colonisation (following
67 Gilbert-Norton et al. 2010), particularly for open-habitat species for which the matrix is
68 impermeable. The use of terrestrial arthropods, particularly carabids and spiders, provides a
69 useful group with which to examine landscape configuration as they are highly speciose,
70 have high reproductive rates, respond quickly to environmental change and contain many

71 different life history stratagems including dispersal, feeding and body size (Duffey 1968,
72 Robinson 1981, Lovei and Sunderland 1996, Marc et al. 1999, Rainio and Niemela 2003).

73 We use this system to examine whether assemblages in transient patches and corridors
74 within the plantation resembled those of the core patches, or instead have been filtered in
75 ways that can be predicted by life history traits. Three alternative colonisation hypotheses
76 are considered for these elements: 1) assemblages in transient stepping-stone and corridor
77 elements resemble those of core patches; 2) assemblages in transient stepping-stone and
78 corridor elements comprise a sub-set of species lacking many specialists restricted to core
79 patches; 3) compared to assemblages within corridors, transient elements have greater
80 representation of dispersive species. We predicted that (P1) assemblages developing in
81 transient patches of open habitat will be represented by small-bodied, aerial-dispersing,
82 habitat generalists, whereas (P2) corridors will have assemblages dominated by large-
83 bodied, active-hunting generalists and that (P3) edge effects will increase the species
84 number and diversity in connectivity elements compared to core patches, due to an influx of
85 vagrant and generalist species.

86 **Material and Methods**

87 Study area and design

88 The study region in eastern England is characterised by a semi-continental climate, sandy,
89 nutrient-poor soil and historically supported extensive semi-natural grass-heath (Dolman
90 and Sutherland 1992). Most grass-heath was converted to arable or forestry during the 20th
91 Century, but remaining grass-heath sites support large numbers of scarce and threatened
92 taxa, characterised by specialist coastal, continental and Mediterranean species dependent
93 on physically disturbed open habitats (Dolman et al. 2012). Most of the remaining grass-
94 heaths are designated for their conservation value but are fragmented and isolated.
95 Regional conservation strategy seeks to improve habitat quality within these remaining
96 grass-heaths, but also to restore functional connectivity between these ‘core patches’, with
97 a focus on connectivity across and within extensive (185 km²) plantation forest (Thetford
98 Forest, 0°40'E, 52°27'N) established over former open habitats in the early 20th Century.
99 During this study Thetford Forest was dominated by conifers, comprising 80% Corsican
100 (*Pinus nigra*) and Scots (*P. sylvestris*) pine, managed by clear-felling (typically at 60-80 years)
101 and replanting of even-aged management patches (hereafter ‘stands’). The forest landscape
102 also contains open-habitat elements that differ in configuration. Narrow linear grass-heath
103 ‘corridors’ are retained along trackways and their perennial verges that form an extensive
104 (1290 km, 18 km²) open-habitat network permeating the forest landscape; following
105 planting of the adjacent tree crop each trackway section provides open conditions for at
106 least 20 years (see below). Larger transient stepping-stone patches (mean area of sampled
107 patches: 9.0 ha ± 8.6 SD) are provided by clear-felled and replanted stands that offer short-
108 lived (5-7 years) open-habitat that is variably connected to the corridor network (Fig. 1);

109 these provide an excellent opportunity to examine colonisation after regular, systematic
110 disturbance events. Vegetation regeneration in these transient patches is derived from
111 persistent grass-heath seed-banks, augmented by zoochorous and aerial dispersal (Eycott et
112 al. 2006a, b, Eycott et al. 2007). Both trackways and transient patches provide fine-grained
113 mosaics of modified grass-heath vegetation and bare mineral soil.

114 Assemblages of 'core patches' were characterised by sampling three locations within each
115 of eight grass-heaths (mean area 106 ha \pm 130 SD), either abutting or located within 2.5 km
116 of the forest (mean=0.72 km \pm 1.11 SD). These included seven Sites of Special Scientific
117 Interest (SSSI) designated for biodiversity under UK conservation legislation, of which five
118 were also designated as Special Areas of Conservation (SAC) under the European Habitats
119 Directive. Most were bordered by plantation forest and or intensive agriculture. Sampled
120 areas comprised physically-disturbed habitats on which characteristic regional biota depend
121 (Dolman et al. 2012), including areas grazed by sheep and rabbits and recently-ploughed
122 areas associated with heathland management.

123 Nineteen transient stepping-stones were sampled. Prior to planting, stands were cleared of
124 coarse woody debris (residue of clear-felling) and planting lines ploughed exposing
125 extensive bare mineral substrate. Bare soil is encroached by ground vegetation within 3-4
126 years (Wright et al. 2009); tree cover then progresses with canopy closure after 20 years
127 (Hemami et al. 2005). Extensive sampling shows open-habitat carabid and spider species
128 persist for approximately five and seven years following planting (Lin et al. 2007, Pedley
129 2012). In the current study sampling considered stands up to seven years (n=5 for one-year-,
130 three-year- and five-year-old; n=4 for seven-year-old). These transient open-habitat patches
131 are fragmented, comprising approximately 6-8% of the 60-80 year growth cycle, while the

132 surrounding plantation crop is impermeable to open-habitat species (Bertoncelj and Dolman
133 2013a, Pedley and Dolman unpublished). Transient patches available for sampling were
134 clustered in the core plantation area (Fig. 1), but distance to the nearest grass-heath site
135 was similar to that of the corridor elements ($t=-1.472$, $DF=30.7$, $P=0.151$; transient
136 mean= $2.1 \text{ km} \pm 1.0 \text{ SD}$, corridor mean= $1.7 \text{ km} \pm 0.6 \text{ SD}$).

137 Fifteen corridor elements were sampled. Trackways (mean width $12.7 \text{ m} \pm 3.9 \text{ SD}$)
138 comprised two elements: a central wheeling with sparse vegetation and exposed substrate,
139 flanked by vegetated verges. Many heathland-associated carabid and spider species have
140 been recorded from this trackway network (Lin et al. 2007, Pedley et al. 2013a) but are
141 excluded by shading from those trackways within or bordering older stands (≥ 20 years)
142 (Pedley et al. 2013a). Sampled verges were each adjacent to thicket-aged stands (11-20 year
143 old, following Hemami et al. (2005) and a mean of $0.78 \text{ km} (\pm 0.25 \text{ SD})$ from the nearest
144 other sampled trackway. To minimise shading effects, for trackways oriented east-west the
145 northern-most verge was sampled, and for those oriented north-south the widest verge was
146 sampled.

147 Arthropod Sampling

148 Ground-active arthropods were sampled in 2009 by pitfall transects that formed the
149 standardised unit of replication for analysis. Although a single sampling year may miss year-
150 on-year differences in compositional detail of sampled assemblages, previous research on
151 ground invertebrates in the study region has shown relatively stable species compositions
152 (Lin et al. 2007, Pedley et al. 2013b). A single pitfall transect was placed in each of the 15
153 corridor elements. Two transects were placed within each of the 19 transient stepping-
154 stones, separated by at least 50 m for independence. In each of the eight heathland sites,

155 three independent transects were again separated by at least 50 m. We acknowledge that
156 replicate transects sampled within each grass-heath site, and within each transient patch,
157 are not spatially independent, and where possible control for this spatial pseudo-replication
158 by including site identity as a random effect in subsequent analyses (see below).

159 Arthropod assemblages were sampled in each transect in May and again in June, the peak
160 periods of arthropod abundance and activity, each time using six pitfall traps (each 7.5 cm
161 deep, 6.5 cm diameter, filled with 50 ml of 70% ethylene glycol), set 15 m apart and opened
162 for seven consecutive days. The two catches were pooled to give one composite sample per
163 transect for subsequent analyses. In August 2009, vegetation height in each transect was
164 assessed at 40 points using a sward stick (diameter 90 mm, weight 250 g, following Dolman
165 and Sutherland 1992), and percentage of bare substrate visually estimated in 20 cm x 20 cm
166 at each point. At each of these points the presence of key structural vegetation features
167 (herbs, grasses, moss, lichen, shrub, and succulent) was also recorded, providing frequency
168 per transect.

169 Carabid identification followed Luff (2007). Adult spiders were identified to species
170 following Roberts (1987, 1996), juveniles and sub-adults with undeveloped reproductive
171 structures were not identified. Analysis of spider assemblages was confined to ground-
172 active species as sedentary invertebrates, such as many web-spinning spider, are poorly
173 represented by pitfall catches (Topping and Sunderland 1992). Ecological and habitat
174 affinities of identified species were classified following Luff (1998, 2007) for carabids and
175 Harvey et al., (2002) and Roberts (1996) for spiders. Habitat affinities comprised: shaded
176 woodland habitats, hereafter 'woodland'; 'generalist' species of multiple or any mesic
177 habitat; and 'dry-open' species associated with dry calcareous or acidic grassland, dry

178 lowland heathland, dunes, sand or gravel pits. Open habitat connectivity elements therefore
179 provide potential habitat for these 'dry-open' species that are excluded from (unable to
180 percolate through) the closed-canopy forest matrix, but their presence depends on
181 colonisation following dispersal.

182 Life history traits

183 Potential ecological factors that may determine assemblage responses to habitat elements
184 were selected. We considering the life history traits of dispersal ability and foraging type
185 (Pedley and Dolman 2014) for which published information was available at the individual
186 species level (Online Resource 1). To ensure ordinal variables had a useable number of
187 species per category, uncommon sub-categories were merged into broader classes. For
188 spiders, aerial dispersal by ballooning (passively floating on silk threads) is considered
189 effective for both short- and long-distance dispersal to colonise suitable habitat (Duffey
190 1998, Bell et al. 2005). Following Lambeets et al. (2008) and Langlands et al. (2011), we
191 assigned spider species as ballooners according to the review by Bell et al. (2005). Carabid
192 dispersal ability was assessed by wing morphology following Barbaro and van Halder (2009),
193 with species exhibiting well-developed wing structures (macropterous) considered effective
194 aerial dispersers, relative to those with rudimentary wing development (brachypterous).
195 Wing-dimorphism (species with both long- and short-winged forms) was classified
196 separately and is considered to provide dispersal advantages over brachypterous species,
197 especially in temporally- and spatially-heterogeneous landscapes (Kotze and O'Hara 2003).

198 Analysis

199 Spiders and carabids were analysed separately, considering composite samples from
200 individual transects. All analyses were carried out using the R statistical software (R
201 Development Core Team 2018). Sampling efficiency was compared between landscape
202 elements using sample-size based rarefaction curves calculated using the iNext package
203 (Hsieh et al. 2016). Assemblage composition was visualised using non-Metric
204 Multidimensional Scaling (NMDS) performed on a matrix of Bray-Curtis dissimilarities of
205 abundance data (square root transformed and Wisconsin double standardization) using the
206 vegan package in R (Oksanen et al. 2018). Habitat structure variables were tested for
207 association with community composition (NMDS) using the envfit function in vegan.

208 Arthropod assemblage composition was compared among core patches, corridors, and
209 transient stepping-stones by multivariate Generalised Linear Models (GLMs) implemented in
210 the mvabund package (Wang et al. 2012) using likelihood-ratio-tests with a negative
211 binomial error distribution.

212 Richness and aggregate abundance of dry-open, generalist and woodland species were
213 compared between landscape elements using Generalised Linear Mixed Models (GLMM).
214 Site (stand, trackway or heath site) was included as a random effect as transects in corridors
215 may capture greater beta diversity than replicates (separated by ≥ 50 m) within individual
216 grass-heaths or transient stepping-stones. Models used Poisson or negative binomial error
217 as appropriate, with landscape element means compared by Tukey pairwise comparisons.
218 GLMMs were implemented in R using the glmer and glmmabmb functions from the
219 packages lme4 (Bates et al. 2015) and glmmADMB.

220 Linkages within trait data (across species) were examined by a Principal Coordinates
221 Analysis (PCoA) applied to a trait-dissimilarity matrix (traits as variables, species as samples),

222 using the Gower coefficient (Gower 1971) using the vegan package. The Gower coefficient
223 was used as it handles mixed data types (nominal, ordinal or continuous – here standardised
224 to zero mean and unit variance). Associations among traits were visualised by plotting trait
225 vectors (from Spearman rank correlation in relation to the first two PCoA ordination axes)
226 following Langlands et al. (2011); species habitat affinities were examined to visualize
227 correlations with species traits. Species were also grouped by family (spiders) or tribe
228 (carabids) to allow a qualitative examination of taxonomic influence in trait space.
229 Phylogenetic character analysis was not possible due to unresolved phylogeny of both the
230 spider and carabid assemblages.

231 Trait-specific responses to habitat types were analysed using fourth-corner analysis (Dray
232 and Legendre 2008) that tests the link between all combinations of species traits and
233 environmental attributes (here: core patch, corridor and transient landscape elements). By
234 re-sampling three data matrices this approach indirectly relates matrix 'R' (environmental
235 attributes x site) to matrix 'Q' (trait x species), via matrix 'L' (species-abundance x site). The
236 fourth-corner procedure uses a generalised statistic S_{RLQ} that is equivalent to the Pearson
237 correlation coefficient r when trait (Q) and environment (R) variables are quantitative, to χ^2
238 when both sets are qualitative, and to the correlation ratio η^2 for mixed data (Dray and
239 Legendre 2008).

240 The observed strength of linkage among traits and environmental attributes is assessed
241 against that which may arise by chance, with null models determined by the observed
242 structure of data matrices, using randomisation and permutation (Dray and Legendre 2008).
243 Permutation Model 1 was used to test the null hypothesis that R is not linked to Q, by
244 examining links between fixed species traits and fixed site characteristics. This model

245 randomises species' relative to site characteristics (permuting within each column of matrix
246 L), but does not re-sample either the species-trait (matrix Q) or environment-site (matrix R)
247 relations; appropriate (Dray and Legendre 2008) as traits were determined from the
248 literature not empirical sampling, while the environment matrix was classified *a priori*. Prior
249 to analysis, arthropod abundance data were square root transformed to reduce the effect of
250 dominant species. We conducted the analysis first on all recorded species, then separately
251 on only those associated with dry-open habitats, and finally for the subset of species
252 recorded on grass-heath sites. Fourth-corner analyses were calculated with 9999
253 permutations, adjusted using false discovery rate correction procedures that account for
254 repeated testing, using the ade4 package (Dray and Dufour 2007).

255

256 **Results**

257 We identified 3660 carabid individuals from 70 species and 12483 spiders from 115 species,
258 of which 11521 spiders from 60 ground-active species were considered in analyses. Pitfall
259 trapping within each landscape element effectively represented carabids and ground-spider
260 with species accumulation curves approaching asymptotes (Online resource 2).

261 Carabid and spider assemblages were also successfully represented in ordination space
262 using NMDS (stress: 0.21 and 0.19 respectively). Both carabid and spider composition (Fig.
263 2) differed significantly between each of the three landscape elements (mvabund GLMs:
264 deviance=758.9, $P < 0.001$; deviance=817.8, $P < 0.001$ respectively; paired comparisons all
265 $P < 0.001$). For carabids, assemblages of core patches were distinct on NMDS axis one, while
266 transient and corridor carabid composition differed from each other on axis two (Fig. 2).
267 Similarly for spider assemblages, axis one separated core patches from both connectivity

268 elements while axis two again separated transient and corridor elements (Fig. 2) though to a
269 lesser extent than for carabids.

270 For both arthropod groups, assemblage composition was significantly associated with all
271 habitat structure variables (Fig. 2, Online Resource 3). Heath sites and assemblages were
272 characterised by greater representation of lichens, succulents, bare ground and short
273 swards; transient patches had taller vegetation, while corridors had greater cover of herbs,
274 shrubs and grasses than transient elements (Fig. 2, Online Resource 3).

275 Arthropod species associated with dry-open habitats occurred across all three landscape
276 elements. For dry-open carabids, mean species richness per sample was similar between
277 core patches and both types of connectivity element, although their aggregate abundance
278 was lower in corridors (Fig. 3, Online Resource 4). Rarefaction showed cumulative richness
279 of dry-open carabids did not differ between core patches, corridors or transient patches
280 (Online Resource 2). For dry-open spiders, mean richness per sample was similar between
281 transient elements and core patches but lower in corridors, while abundance was lower in
282 transient patches than core patches and was again lowest in corridors. Cumulative richness
283 of specialist spiders was also significantly lower in corridors (Online Resource 2). Whilst core
284 patches were dominated by dry-open species, both types of connectivity elements had
285 greater representation of generalist and woodland species. For generalists, mean
286 abundance and richness per sample were similar between transient and corridor elements
287 for both taxa, with generalists particularly dominant in spider assemblages (Fig. 3).

288 Rarefaction showed similar rates of species accumulation of generalists across elements for
289 both taxa (Online Resource 2). Species associated with woodland were particularly scarce in
290 the open grass-heath sites (with only two woodland spider individuals and 33 woodland

291 carabid individuals recorded), but contributed to assemblages in both connectivity
292 elements, with woodland spiders having greater mean abundance and richness in corridors
293 than transient patches (Fig. 3).

294 Inter-relationship among traits

295 The first two axes of the PCoA represented 57.9% and 60.8% of the variation in traits among
296 carabid and spider species respectively (Fig. 4, Online Resource 5), indicating inter-
297 correlation among traits in each case. For carabids, PCoA axis one was positively associated
298 with larger body size, brachypterous wing morphology and carnivorous feeding, and
299 negatively associated with inter-correlated traits of herbivory and macroptery. Carabids
300 associated with dry-open habitat were negatively related with PCoA axis one and positively
301 with axis two, habitat generalists were negatively related with axis two. Distribution of
302 carabid tribes within the PCoA suggested some phylogenetic pattern, with herbivory only
303 found in the Zabrinini (*Amara* and *Curtonotus*) and Harpalini (*Harpalus*, *Bradycellus*,
304 *Ophonus*), which tended to be macropterous and mainly associated with dry-open habitats.
305 Wing morphology was not strongly phylogenetically conserved in remaining carabid tribes.

306 For spiders, PCoA axis one was positively associated with ambush hunting and habitat
307 generalists and negatively associated with dry-open habitat association and a running
308 hunting strategy. Spider PCoA axis two was positively associated with strongly inter-
309 correlated traits of running hunting strategy and larger body size, and with flight dispersal,
310 and was negatively associated with woodland habitat association and a stalking hunting
311 strategy. The spider PCoA showed clustering of the main families (Fig. 4). The large, running
312 hunters of the families Lycosidae, Gnaphosidae and Clubionidae were grouped towards the
313 upper part of the ordination, but varied along axis one between species associated with dry-

314 open habitats (in the upper-left of the ordination) and generalist species (upper right).
315 Thomisidae, Theridiidae and Salticidae, were associated with ambush and stalking hunting
316 strategies (lower part of the ordination) and tended to be smaller-bodied and habitat
317 generalist or woodland-associated species.

318 Species traits among landscape elements

319 Fourth corner analysis revealed substantial differences in the trait composition of
320 assemblages between the three landscape elements. Of the seven carabid and five spider
321 traits tested, most correlated significantly with at least one landscape element (Table 1).
322 Notably, all traits that significantly correlated with transient and corridor elements showed
323 an opposite direction of correlation between these connectivity element types. Both carabid
324 and spider assemblages in transient patches were correlated significantly with larger body
325 size. Transient patches were also positively correlated with a carnivorous diet for carabids
326 and with a running-hunting strategy for spiders. Importantly, traits conveying an increasing
327 dispersal ability by flight (carabid macroptery) or aerial drift (spider ballooning) were not
328 positively correlated with either type of connectivity element. For carabids, macroptery was
329 negatively correlated with corridors, while for spiders ballooning was negatively correlated
330 with transient elements. Wing-dimorphism for carabids was positively correlated with
331 corridors and core patches, but negatively correlated with transient elements. Trait
332 correlations tended to be reversed between core patches and transient elements; for
333 example, small-bodied herbivorous carabids and small-bodied ambush-hunting spiders were
334 correlated with core patches.

335 When considering only those species associated with dry-open habitats, several important
336 trait-habitat correlations were apparent. For dry-open carabids, corridors were correlated

337 with a carnivorous diet and lack of flight dispersal (brachyptery), while transient elements
338 were correlated with large-bodied, herbivores with flight dispersal (Table 1). For dry-open
339 habitat spiders, transient elements were again correlated with large, running hunters
340 lacking ballooning dispersal, while corridors were not correlated with particular traits.
341 Finally, restricting fourth-corner procedures to only species recorded in core patches did not
342 provide any addition correlations for either taxa (Online Resource 6) with results largely
343 consistent to previous analyses of the entire assemblage.

344 **Discussion**

345 Ground-active carabid and spider assemblages in core patches, transient stepping-stones
346 and corridor connectivity elements differed in both species composition and trait
347 representation, suggesting filtering by traits. Consequently connectivity elements differed in
348 functional composition, both from each other and from the remaining grass heaths, but
349 nevertheless increased beta diversity at the landscape-scale and supported obligate dry-
350 open habitat species characteristic of core patches of grass heath. While corridors contained
351 far fewer dry-open individuals than core patches, transient stepping-stone elements had
352 similar mean dry-open richness to that in core patches. Our data on trait filtering did not
353 resolve to generalised colonisation hypotheses that cut across these two taxa, due to
354 differences between beetles and spiders in dispersal mechanisms and also their patterns of
355 associations among traits. However, patterns regarding habitat association, generalist
356 domination of connectivity elements, and increased species diversity in connectivity
357 elements were consistent with our predictions (P3), while dispersal, habitat structure (patch
358 quality) and edge effects (patch configuration) all appeared to influence differences in trait
359 composition between elements.

360 **Assemblage composition among landscape elements**

361 We predicted that generalist species would dominate the connectivity elements and that
362 richness and abundance would be greater in these patches, as expected from edge- and
363 ecotone-effects. We found that generalist species made up a large proportion of the carabid
364 assemblage and dominated spider assemblages, in both types of connectivity elements,
365 with greater overall species richness of both taxa than in core patches of grass heath
366 habitat. While both types of connectivity element also contained significant numbers of

367 specialist carabids, core patches contained few species that were not specialists of dry-open
368 habitat. Our results are consistent with others that find generalist species resilient to
369 landscape change and dominant in disturbed or human-altered landscapes (Robinson and
370 Sutherland 2002, Ewers and Didham 2008, Smith et al. 2015). As generalist species tend to
371 be those associated with invasions and also better able to cope with changing, fragmented
372 and disturbed ecosystems (Marvier et al. 2004), large numbers of generalists may be
373 considered a concern for the conservation of natural and semi-natural assemblages.

374 Compositional edge effects are well known for forest boundaries (Downie et al. 1996, Muff
375 et al. 2009, Campbell et al. 2011, Kowal and Cartar 2012); Ewers and Didham (2008) found
376 increased beetle diversity at forest/grassland boundaries with compositional differences
377 extending over 1 km. The greater richness of generalist and woodland-associated species
378 within our corridor and transient elements, than in core patches, may partly be attributable
379 to an influx of 'vagrants' from the adjacent matrix. Inflated species richness in narrow and
380 small habitat patches is well documented for invertebrates (Halme and Niemela 1993,
381 Driscoll and Weir 2005). Webb and Hopkins (1984) found that small heathland fragments
382 supported greater invertebrate richness than larger intact sites, while Halme and Niemela
383 (1993) found similar patterns in carabids inhabiting small forest fragments in Finland. In the
384 current study it is likely that the narrow shape of corridors and the relatively small size of
385 transient elements (and hence large edge ratios), favoured the incursion of woodland and
386 generalist species from the matrix.

387 However, the greater species richness in connectivity elements also reflects the greater
388 structural heterogeneity of habitats within these elements (Bieringer et al. 2013), not just
389 edge-incursion ('spill-over') from the matrix. Both types of connectivity element contained a

390 juxtaposition of: open micro-sites with bare disturbed mineral soil (trackway wheelings and
391 ploughed planting rows), taller grassy herbaceous vegetation (trackway verges and baulks
392 between planting rows) and woody ecotones at their margins, providing opportunities for
393 species associated with these individual habitats but also those requiring micro-habitat
394 juxtaposition and mosaics (Dolman et al. 2012). Similarly, Driscoll and Weir (2005) found
395 linear strips of Australian mallee habitat were species-rich due to both 'strip-specialists' and
396 matrix species. Assemblages in the core patches of grass heath habitat may represent not so
397 much a pristine reference sample, but a sub-set of the landscape-wide biota filtered to
398 exclude generalist and woodland associated species. This reflects the historic development
399 of heathland from degraded pasture-woodland (Fuller et al. 2017).

400 Importantly, we found that the ground-active carabid and spider assemblages in
401 connectivity elements were significantly different to each other as well as to assemblages of
402 core patches. Although generalist abundance and richness did not differ between transient
403 stepping-stones and corridor elements for either arthropod group, the mean richness and
404 abundance of specialist dry-open associated arthropods was greater in transient stepping-
405 stone elements than in corridors. For spiders, both mean (per sample) and cumulative
406 (measured through rarefaction) specialist richness were similar between transient elements
407 and core patches, but significantly lower in corridors. This is despite an expectation of
408 efficient dispersal in linear elements with resistant (i.e. internally reflecting) borders set in a
409 'hard' matrix (i.e. one that is unlikely to provide dispersal or breeding habitat for target
410 species) (Baum et al. 2004, Ockinger and Smith 2008, Bertonecelj and Dolman 2013b). This is
411 also contrary to findings for a Hemipteran species, where in a soft (low-resistance) matrix,
412 both corridors and stepping stones significantly improved connectivity, but in a hard matrix

413 stepping-stone elements did not improve dispersal over non-connected controls (Baum et
414 al. 2004). Despite the expectation of greater ease of colonisation along linear elements in a
415 non-permeable matrix, other mechanisms may contribute to the greater abundance of
416 specialist dry-open associated species in transient stepping-stones. First, although corridors
417 are often conceptualised as conduits for individual movement and colonisation between
418 habitat patches, diffuse resident populations may breed and disperse by percolation over
419 several generations. Second, the larger transient patches, with reduced shading by adjacent
420 forest and extensive disturbed mineral soil, may have provided greater habitat suitability for
421 specialists than provided by corridors. Third, the larger size of transient patches relative to
422 narrow corridors may have facilitated their colonisation by aerial-dispersing species.

423 Trait filtering

424 We predicted that assemblages developing in transient patches of open habitat would be
425 represented by small-bodied, aerial-dispersing species, whereas corridors would be
426 dominated by large-bodied, active-hunters. Trait correlations were in fact opposite to this
427 prediction. However, while small body size and traits conveying greater dispersal ability
428 were not more prevalent in transient patches overall; when analyses were confined to
429 specialists of dry-open habitat (that are excluded from and do not percolate through the
430 matrix), flight dispersal significantly correlated with transient elements for carabids. Where
431 a significant trait correlation was detected with both corridor and transient elements, these
432 were in opposing directions, indicating differential filtering of assemblages in these
433 connectivity elements.

434 Differences in dispersal trait correlations were evident between the two arthropod groups
435 and no consistent functional response to either connectivity type was found. The ability to

436 disperse aerially is seen as a key trait for insects colonising unstable habitat (Roff 1990,
437 Ribera et al. 2001) and ballooning spiders are often considered early colonists to newly-
438 opened or disturbed habitat, such as after volcanic eruptions (Crawford et al. 1995) or in
439 agricultural fields (Nyffeler and Sunderland 2003, Schmidt and Tschardt 2005). Prevalence
440 of aerial dispersal was only significantly greater in core patches for spiders and in transient
441 patches for dry-open carabids. The findings for dry-open carabids are consistent with results
442 in other systems (Gobbi et al. 2007, Moretti and Legg 2009, Wamser et al. 2012). Moretti
443 and Leg (2009) reported large, highly-mobile insects respond to disturbance caused by
444 regular winter fires, while Gobbi et al. (2007) found that younger sites with less stable soil
445 conditions favouring greater representation of winged carabids in an alpine glacial
446 chronosequence. In our long-lived corridors, dry-open carabids consisted of flightless
447 carnivorous carabids, consistent with results from other stable or linear connectivity
448 elements (Ribera et al. 2001, Wamser et al. 2012).

449 While disturbed and early-successional habitats have greater representation of flight
450 capable species (Gutierrez and Menendez 1997, Ribera et al. 2001, Pedley and Dolman
451 2014), landscape configuration may also influence species distributions at the regional scale.
452 Negative correlation of ballooning capable spiders with transient stepping-stone elements
453 may in part be due to fragmentation of suitable habitat; with passive dispersal representing
454 a high-risk strategy for narrow-niche species in fragmented landscapes. Bonte et al. (2003b)
455 showed that specialist xerophilic spiders in fragmented sand dune habitat were less likely to
456 balloon than habitat generalists. We found generalists were associated with flight dispersal
457 for spiders but not for carabids, although brachyptery was associated with woodland

458 carabid species, as expected by affinity of flightless species to more stable habitats (Ribera
459 et al. 2001, Pedley and Dolman 2014).

460 For local dispersal, terrestrial movement alone may be sufficient for many ground-active
461 species (Samu et al. 2003, Brouwers and Newton 2009). In the current study, large carabids
462 and spiders were correlated with transient stepping-stone elements but not corridors, even
463 when analysis was restricted to dry-open specialists. Results of a systematic review by
464 Brouwers and Newton (2009) showed that larger carabids covered more ground per day
465 than small species, likely due to both greater food requirements (Lovei and Sunderland
466 1996) and greater movement capability. Kormann et al. (2015) also found larger arthropods
467 tended to benefit more from increased landscape connectivity, in the form of more closely-
468 clustered grassland fragments, than did smaller species. For spiders, larger body size was
469 associated with running hunters, of which Lycosidae made up the majority of the sampled
470 assemblage. These running hunters have potential terrestrial dispersal distances of several
471 hundred metres over a lifetime, with daily ranges recorded between 2-50m (Kiss and Samu
472 2000, Bonte et al. 2003a, Bonte et al. 2007). Such lifetime distances combined with high
473 daily activity rates may explain the positive correlation of running hunters with transient
474 patches, compared to less active hunting strategies (ambush and stalking) that are
475 associated with smaller body size in spiders.

476 Conclusions

477 Concepts introduced through the theory of island biogeography and metapopulation
478 dynamics suggest increasing habitat fragmentation will restrict species with poor dispersal
479 to persistent refuges, while those with enhanced dispersal ability can take advantage of
480 transient and connecting habitat elements. In the plantation landscape studied here, we

481 found that the assemblages in core patches, corridors and transient stepping-stones were
482 dissimilar in terms of both species' composition and trait representation. Nevertheless,
483 transient elements supported similar richness of characteristic dry-open associated
484 specialists for which remnant heaths are designated. Dry-open habitat species in these
485 transient patches were predominately large running spiders and large herbivorous carabids.
486 However, dispersal strategies for these two arthropod groups were likely very different. In
487 contrast, narrow linear corridors were less favourable to the open heathland assemblage,
488 with spiders dominated by generalists. To provide connectivity for less-mobile species and
489 taxa, corridors should reduce edge-related effects, as wider higher-quality corridors are
490 likely to support more specialist species (Lees and Peres 2008, Pedley et al. 2013a).
491 Carabids' ability for directional flight, compared to spiders' passive ballooning, may explain
492 their greater ability to take advantage of transient connectivity elements within complex
493 landscapes. Contrasts between carabids and spiders in their response to landscape
494 configuration, emphasise the need for assessments to examine multiple taxa before making
495 generalisations. Dispersal corridors and transient stepping-stones provided different trait
496 and functional composition and were not interchangeable in this system.

497 Fig. 1. The distribution of core patches of grass-heath, corridor and transient stepping-stone
498 elements sampled within the study landscape.

499 Fig. 2. Non-metric multi-dimensional scaling (NMDS) ordination of ground-active carabid
500 and spider assemblages (stress = 0.21 and 0.19 respectively). Open circles represent
501 sampled transects and shaded ellipses the 95% confidence intervals of habitat element
502 centroids. Significant habitat structure variables are displayed as vectors, longer vectors
503 represent stronger association with species ordination. Online Resource 3 gives the vector,
504 R^2 and p-value for each environmental variable.

505 Fig. 3. Mean (+/-se) of ground-active carabid and spider abundance and species richness
506 shown separately for generalist, woodland-associated and dry-open habitat specialists,
507 across three landscape elements. Means are calculated from standardized pitfall trap
508 transects where each sample represents a composite of two sampling periods. Means that
509 share a superscript (homogenous sub-sets, a-c) do not differ significantly (Tukey pairwise
510 comparisons, $P > 0.05$). Online Resource 4 gives model statistics, p-values and means for
511 each comparison.

512 Fig. 4. Principal coordinate analysis (PCoA) showing associations among ten traits of 69
513 carabid species and eight traits of 59 spider species. Trait vectors represent the Spearman
514 correlations, with the length and direction indicating the relationship with composite PCoA
515 axes (see Online Resource 1 for trait details and Online Resource 5 for trait loadings).

516

517 Table 1. Fourth-corner test results in which different landscape elements are related to
 518 species traits (see Online Resource 1 for trait details). Fourth-corner analysis used
 519 permutation Model 1 and 9999 permutations (see Online Resource 6 for model statistics).

	Carabid - all species			Carabid – dry-open		
	Transient	Corridor	Core patch	Transient	Corridor	Core patch
Body size	+		-	+		-
Carnivorous	+		-		+	-
Herbivorous		-	+	+	-	
Omnivorous						
Macropterous		-		+	-	-
Brachypterous			-	-	+	+
Wing-dimorphic	-	+	+			
	Spiders - all species			Spiders – dry-open		
	Transient	Corridor	Core patch	Transient	Corridor	Core patch
Body size	+		-	+		-
Running hunter	+		-	+		-
Ambush hunter	-		+			
Stalking hunter						
Flight (ballooning)	-		+	-		+

520

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763 **Electronic Supplementary Material:**

764 Online Resource 1: Description of species traits

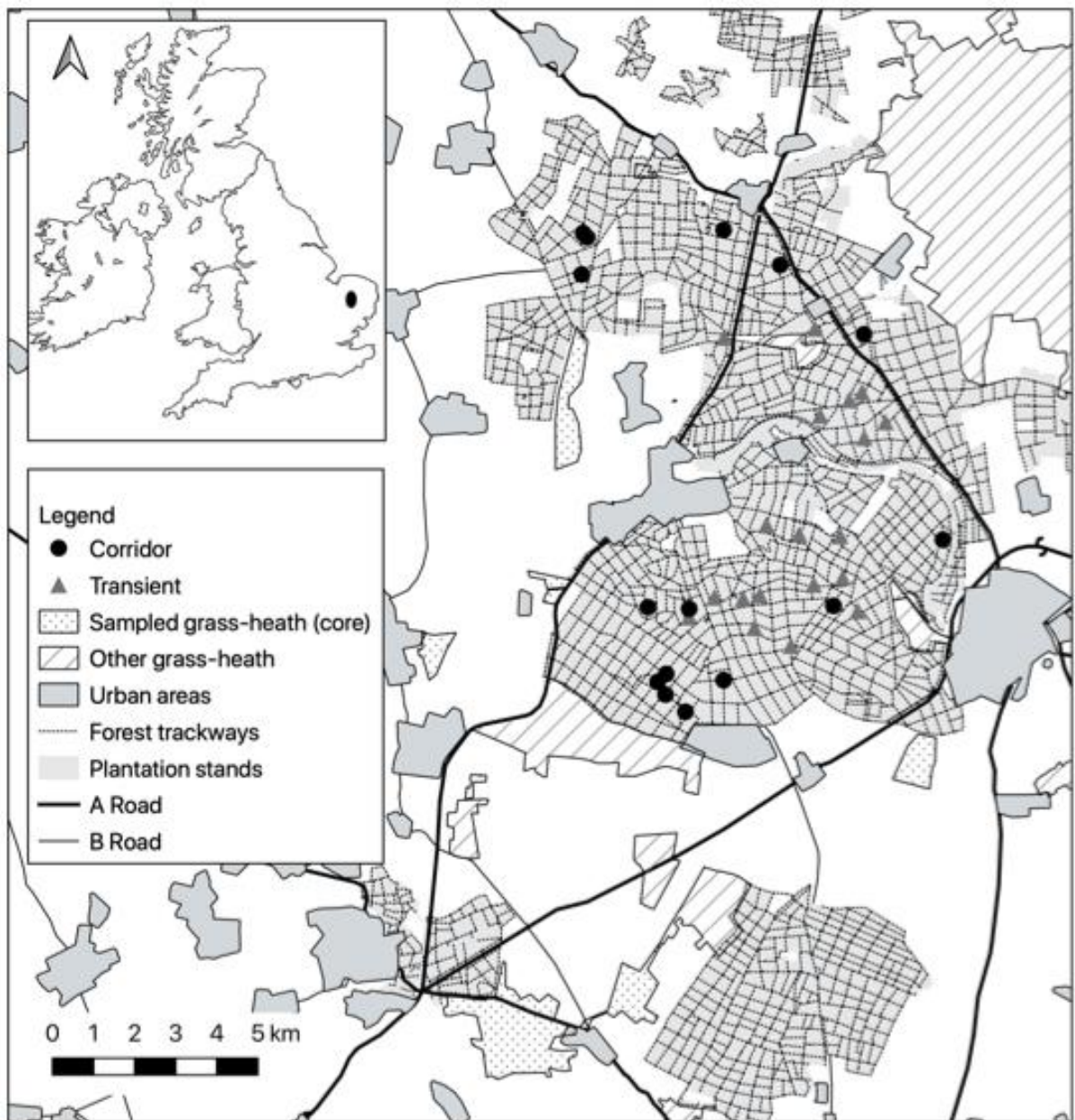
765 Online Resource 2: Rarefaction curves for the sampled arthropod communities

766 Online Resource 3: Summary of habitat structure variables and vector loadings

767 Online Resource 4: Results of Mixed Models

768 Online Resource 5: PCoA loadings

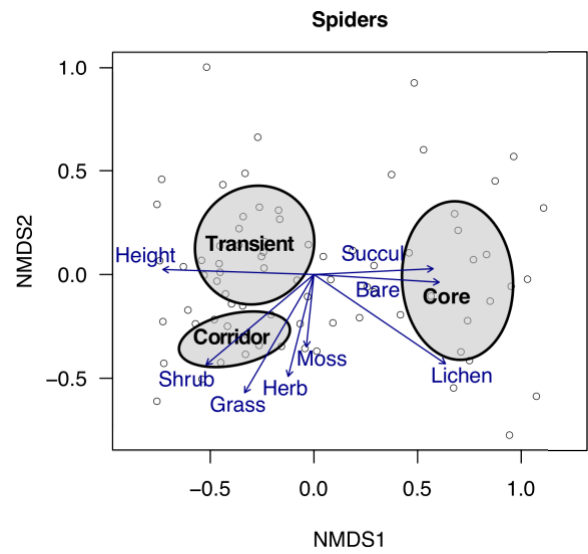
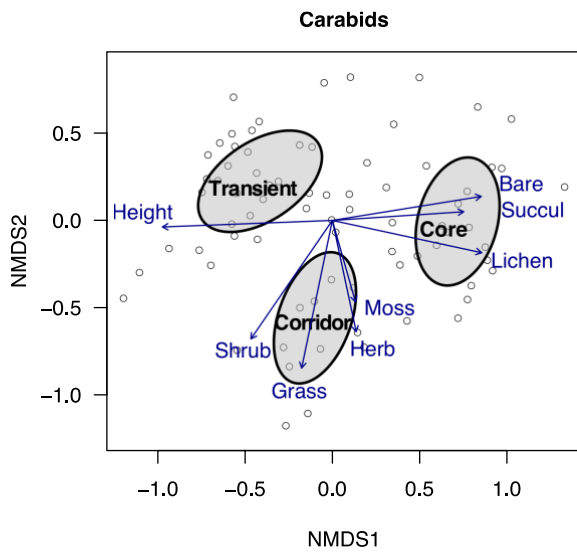
769 Online Resource 6: Fourth-corner model statistics



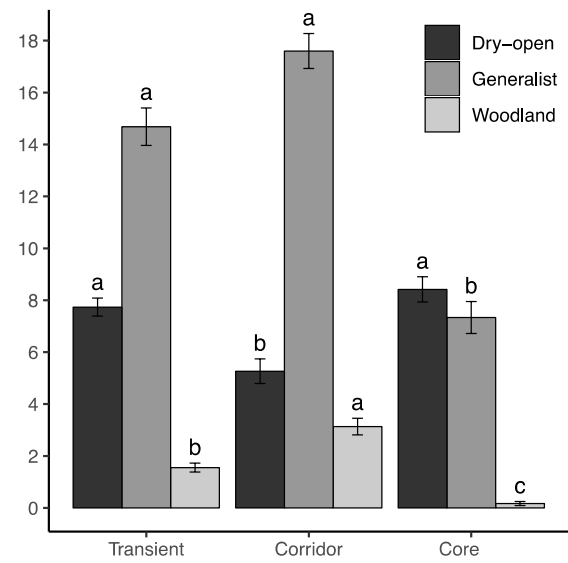
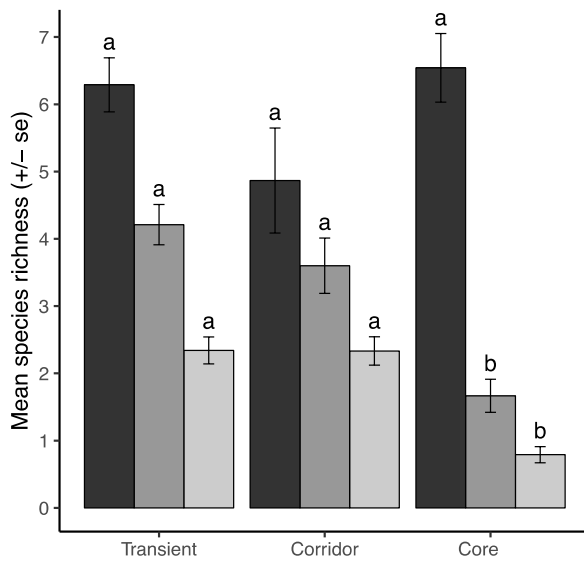
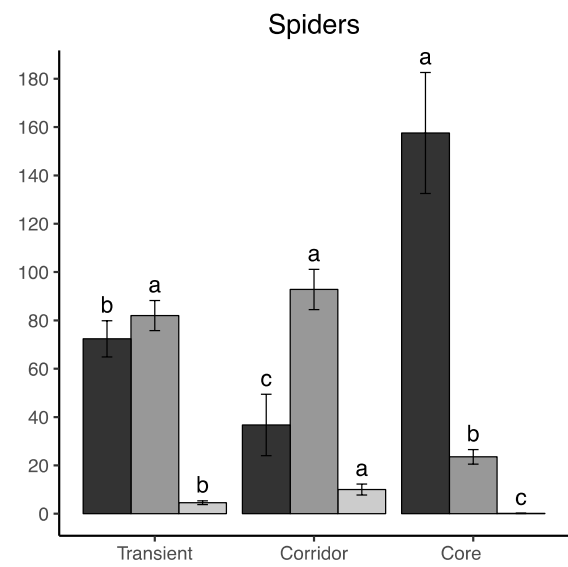
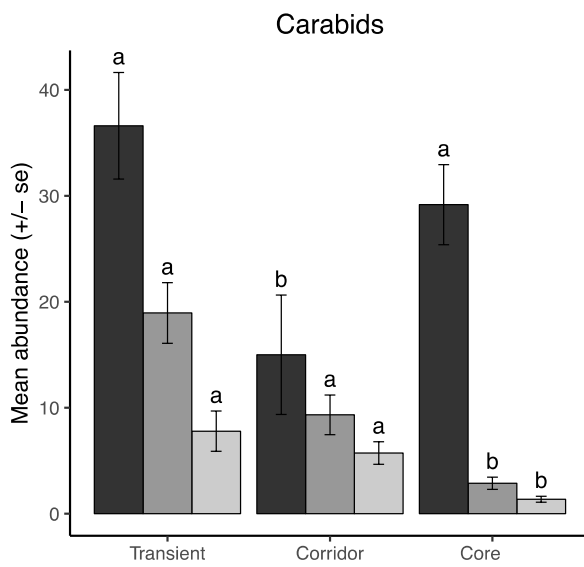
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