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Contrasting beetle assemblage responses to cultivated farmlands and native woodlands in a dynamic agricultural landscape

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Abstract. There is an urgent need to identify ways of managing agricultural landscapes for biodiversity conservation without reducing food production. Farming practices that consider spatio-temporal heterogeneity of farm fields may be a feasible alternative to large-scale revegetation of farmlands for maintaining arthropod biodiversity and their important ecological function. We examined seasonal differences in beetle assemblages in woodland remnants and four adjoining farmland uses in a highly modified agricultural landscape in southeastern Australia. The farmland uses were crops, fallows, and two restoration treatments (fine woody debris applied over harvested crop fields, and restoration plantings). Unexpectedly, overall species richness was significantly lower in remnants than in adjacent farmlands. Remnants and farmlands supported significantly different assemblages, with a third of species found in both habitats. Abundance responses were taxon-specific and influenced by interactions between land use and season. In particular, predator abundance was significantly higher in plantings and fallows during spring compared to summer. Detritivore abundance was significantly higher in the woody debris compared to the adjacent remnants. Herbivore abundance did not differ between remnants and farmlands over time. Complex responses provide strong support for a mosaic of land uses to effectively conserve different beetle groups. Species richness results suggest that further agricultural intensification, in farm fields and through the removal of remnant vegetation, risks reducing beetle diversity in this region. Maintaining farmland heterogeneity with a mix of low-intensity land uses, such as conservation tillage, crop–fallow rotation, restoration plantings, and the novel application of fine woody debris over cultivated fields, may provide seasonal refuge and resources for beetles.

Key words: Coleoptera; fragmentation; landscape mosaic; matrix; restoration; tillage.

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INTRODUCTION

Agriculture is a major cause of decline in biodiversity and ecosystem services globally (Bradshaw 2012, Newbold et al. 2015, Soliveres et al. 2016), due to widespread conversion and degradation of

natural habitats (Gibson et al. 2011, Tilman et al. 2011) and increased intensification of agricultural practices (Benton et al. 2003, Hendrickx et al. 2007, Attwood et al. 2008). Retaining and restoring native vegetation has been identified as critical for conserving biodiversity in agricultural landscapes.

This is because higher species richness and abundance of many taxonomic groups are typically found in semi-natural habitats than in intensive land uses due to higher habitat heterogeneity and resource and niche availability associated with undisturbed, natural vegetation (Benton et al. 2003, Fischer et al. 2006, Attwood et al. 2008). Revegetating areas with native trees and shrubs in highly simplified landscapes also can improve biodiversity by providing habitat and increasing connectivity for some taxonomic groups (Gibb and Cunningham 2010, Lindenmayer et al. 2010, 2016).

While biodiversity conservation has traditionally focused on species associated with patches of remnant native vegetation, there has been growing emphasis on the effect of spatial heterogeneity of the overall landscape on species distribution and assemblages (Fahrig and Nettle 2005, Turner 2005, Vasseur et al. 2013). This has led to repeated calls for conservation ecologists to shift away from a binary patch/matrix perspective to a mosaic view of varying land uses (Ricketts et al. 2001, Bennett et al. 2006, Vasseur et al. 2013), especially in human-dominated landscapes where little or no natural habitat remains (Pimentel et al. 1992, Fournier and Loreau 2001, Bradshaw 2012). More recently, intensively managed farm fields—such as crop monocultures, which typically form the bulk of agricultural landscapes—have been suggested as important drivers of population dynamics and persistence due to their “hidden” habitat value in space and time (Vasseur et al. 2013). This hidden heterogeneity refers to the diversity in management practices and crop types, as well as more subtle changes in crop fields within and between growing seasons. For example, short-term disturbances such as agrochemical use, crop harvest, and grazing, as well as crop rotation over longer time periods (Baudry and Papyz 2001, Bennett et al. 2006), can influence resource availability and affect meta-population dynamics at different spatio-temporal scales (Burel and Baudry 2005, Holland et al. 2005, Bennett et al. 2006). The effects of fine-grained spatio-temporal changes of farm fields on the structure of biotic communities, however, have been rarely studied (Gagic et al. 2012, Vasseur et al. 2013, Puech et al. 2015).

Few studies have concurrently examined a wide range of farm and restoration management options for improving biodiversity (Scott and Anderson 2003, Bridle et al. 2009, Vasseur et al. 2013), while

at the same time taking into account the hidden heterogeneity within farm fields (Vasseur et al. 2013). Management changes to alter the mosaic of resources within farm fields may present more cost-effective and practical options for increasing food production (Pywell et al. 2015) while maintaining biodiversity and ecosystem function (Benton et al. 2003, Tschardt et al. 2005). This is important because not only does large-scale revegetation of productive farmlands reduce food production, but there are large knowledge gaps in the effectiveness of revegetation (especially for arthropod communities, Hunter 2002, Barton and Moir 2015) compared with other farm management strategies (Tschardt et al. 2005, Holzschuh et al. 2009) potentially compatible with sustainable agricultural intensification (Pywell et al. 2015). For example, farming practices that increase landscape heterogeneity, such as applying fallow rotation and other measures to increase groundcover structural complexity, may be as effective as revegetation in improving farmland biodiversity for some taxonomic groups (Benton et al. 2003).

Arthropods comprise a major component of terrestrial biodiversity and provide important ecological functions in agricultural landscapes such as biological pest control (Lovei and Sunderland 1996, Kromp 1999), pollination, decomposition, and weed control (Grimbacher et al. 2006). However, limited knowledge of the distribution and ecology of many species—including habitat requirements in modified landscapes—impedes their conservation and the maintenance of the ecosystem services they provide (Holland et al. 2005, Cardoso et al. 2011, Marrec et al. 2015). Beetles (Coleoptera) are an ideal group for studying impacts of landscape modification because they are speciose and represent a wide range of trophic and functional groups (Lawrence et al. 2000). Beetles are also expected to respond to management actions because they are sensitive to small-scale changes in habitat and seasonal conditions (Bromham et al. 1999, Gibb and Cunningham 2010, Woodcock et al. 2010).

In agricultural landscapes, more structurally complex habitats (e.g., woodlands, plantings, and fallows) generally support higher species richness and abundance of all trophic groups of beetles and more specialized subfamilies of beetles compared to intensive land uses (e.g., crops; Lassau et al. 2005, Attwood et al. 2008, Newbold et al. 2015). Responses of different beetle groups

might also fluctuate over time due to seasonal changes in food availability, habitat quality, or species life cycle (Janzen 1973, Thiele 1977, Grimbacher and Stork 2009). For example, species richness and abundance of most beetle groups might decline between spring and summer due to drier conditions in summer (Hill 1993). However, stronger declines are more likely in intensively cropped land uses due to removal of resources during summer harvest (Sackmann and Flores 2009).

Here, we compared ground-dwelling beetle assemblages between woodland remnants and four adjoining farmland uses comprising crop, fallow, and two restoration treatments (fine

woody debris applied over harvested crop fields and restoration plantings). Our key research question was as follows: What are the differences in beetle assemblages between woodland remnants and adjacent farmlands, and over a crop-growing season (spring and summer)?

MATERIALS AND METHODS

Study site and sampling design

Our study area was a highly modified mixed-cropping landscape within the Lachlan River catchment, New South Wales, southeastern Australia (Fig. 1). Widespread clearing for agriculture has restricted native *Eucalyptus* woodland remnants to

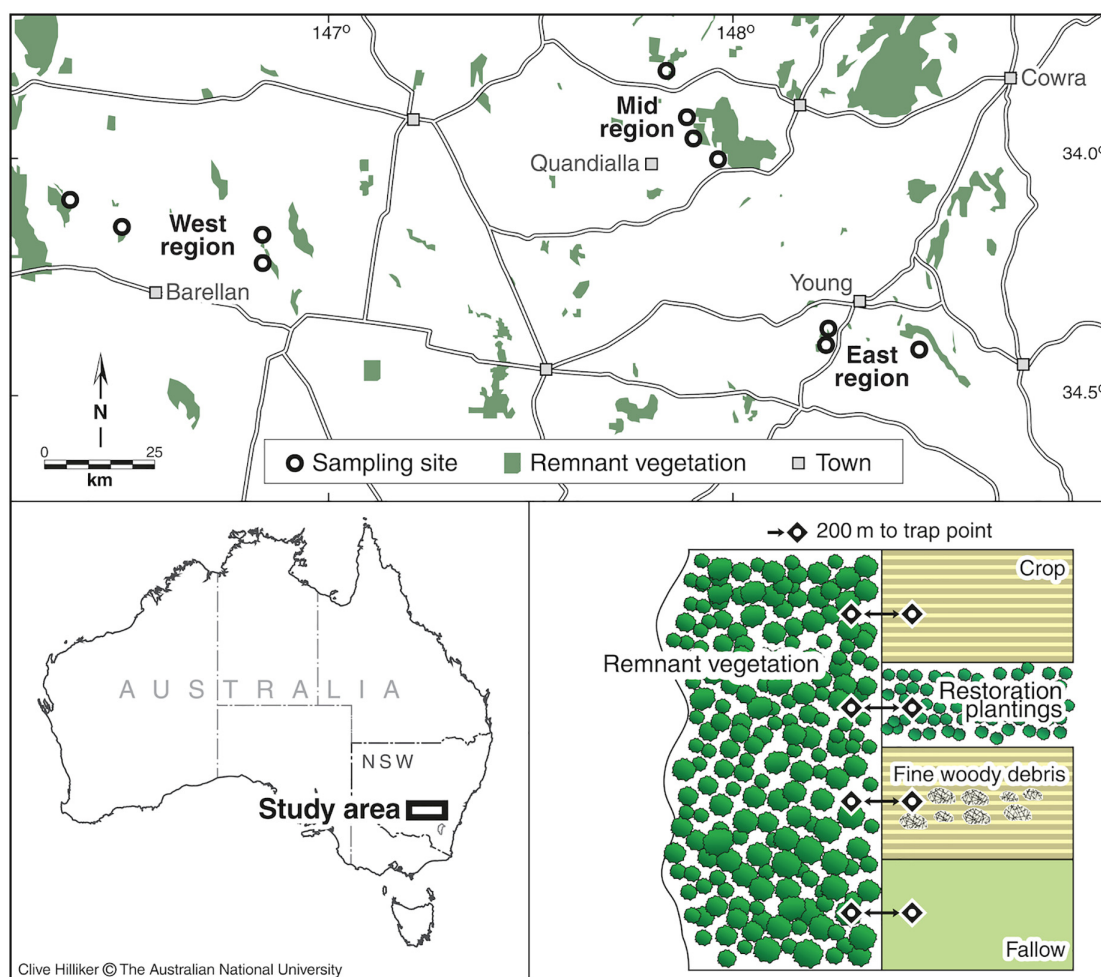


Fig. 1. Map showing study sites in New South Wales, southeastern Australia. Inset shows stylized image of experimental design and pitfall trap placement along four 400-m transects between a remnant patch habitat and adjoining farmland habitats.

infertile steeper areas (Hitchcock 1984, Bradshaw 2012), with many remnants modified by livestock grazing, weed invasion, and changed fire regimes (Norris and Thomas 1991). Our study sites were clustered in three regions along a decreasing elevation and rainfall gradient from the east, mid to west (Fig. 1).

For the purposes of our study, we defined a “patch” as remnant woodland vegetation surrounded by a mostly cleared farmland “matrix”. We focused on eleven remnant patches [patch size $4010 \text{ ha} \pm 486.4 \text{ ha}$ (mean \pm standard error)] selected on the basis that they were *Eucalyptus* woodland communities with high ground-, mid-, and over-story native vegetation complexity (Appendix S1: Fig. S1), with the following adjoining farmland types: (1) winter wheat crop, (2) fallow (rested from crop rotation or sown-pasture rotation), (3) restoration plantings of native shrubs and trees (<7 yr old), and (4) application of eucalypt-based fine woody debris over wheat stubble after harvest prior to sampling (January 2015, Fig. 1; Appendix S1: Fig. S2). The fine woody debris addition is a novel treatment not used previously and piloted in our study to increase ground layer complexity in crop fields to provide resources for arthropod biodiversity. Farmers in our study area were receptive to this treatment because it does not impede cropping machinery unlike larger, coarse woody debris (logs) previously used to restore structural complexity in pastoral areas in Australia (Manning et al. 2013).

Beetle sampling

We used a split-plot sampling design where each remnant patch was matched with the four different farmland matrix types (Fig. 1). We sampled beetles along a 400-m transect from 200 m in each patch out into 200 m in each of the four adjoining farmland matrix types. For consistency in terminology, we referred to each of the four matched patch–matrix combination as a “transect”, and referred to either the matrix (which aggregated four matrix types) or patch side as “habitat.” We then sampled beetles with a pair of pitfall traps located at each end of the transect: 200 m inside the remnant patch and 200 m in the adjoining farmland matrix (Fig. 1). We chose 200 m because it represented the interior position in smaller farm fields. Individual traps from each

pair were placed on either side of a drift fence (60 cm long \times 10 cm high) to help direct arthropods into the trap. Traps were plastic jars (6.5 cm diameter, 250 mL) dug into the ground with the rim level with the soil surface, filled with 100 mL of preservative (1:3 glycol–water mixture, and a drop of detergent to reduce surface tension).

We sampled from the same pitfall trap locations during two distinct periods of the cropping cycle (referred to as “time” in our study): spring when crops were at peak flowering, and summer after crop harvest (stubble retained). A total of 88 pairs of traps (11 replicate sites \times 4 transects \times 2 trap pairs) were opened for 14 d during spring (October–November 2014) and summer (January–February 2015).

Arthropods were preserved in 70% ethanol. All adult beetles were removed and sorted to family and to genus or species where possible. Beetle taxonomy followed Lawrence and Britton (1994) and Lawrence et al. (2000). Where specimens could not be identified to genus or lower, measures of abundance and richness corresponded to morphospecies (sensu Oliver and Beattie 1996), henceforth referred to as species. Each species was assigned to one of three generalized trophic groups: predators, herbivores, and detritivores (including fungivores), based on the predominant feeding behavior of adults at the family and subfamily level, where possible (Lawrence and Britton 1994). We assigned all carabids as predators because purely phytophagous species are considered uncommon in Australia (Gibb et al. 2017). We acknowledge that aggregating data by trophic and family groups may conceal species-level variation. However, this approach is an acceptable compromise for estimating species richness in highly diverse regions where taxa are still poorly described (Ricketts et al. 2001). Higher-level family- and trophic-level patterns may also help provide some capacity to generalize responses for functional groups and infer broad ecological processes (Tscharntke et al. 2005).

Statistical analyses

Samples from each pitfall trap pair were pooled to provide one sample per trap point. Traps at ten sampling points were damaged by vertebrate fauna and discarded from analysis, leaving 166 trap points in total.

We first examined differences in beetle species composition between remnant patches and four farmland matrix types using permutational multivariate analysis of variance (PERMANOVA), based on Bray–Curtis dissimilarities. We ran 999 permutations, and stratified within site and transect to account for the nested sampling design of transects within sites. Singleton species were excluded, and we ran comparisons for spring and summer separately. *P* values were adjusted using sequential Bonferroni corrections to account for multiple comparisons. We used the “vegan” R package for PERMANOVAs (Oksanen et al. 2013).

To identify whether farmland use had an important effect on beetle assemblages in the remnant patch and/or farmland matrix and any interactive effects with time, we used generalized linear mixed-effect models (GLMM, Bolker et al. 2009). Response variables analyzed were the species richness and abundance of the overall assemblage, trophic groups, and the 15 most common families (see Appendix S1: Table S1). The main fixed effects tested were the two-way interactions of transect (four levels: planting, fallow, crop, and woody debris) and habitat (two levels: patch and matrix) or the three-way interactions of transect, habitat, and time (two levels: spring and summer). We controlled for possible effects of region (three levels: east, mid, and west) and remnant patch size (continuous variable) by including them as additive fixed effects, although these factors were not of primary interest in this study. We fitted site, transect location, and trap location as nested random effects (1|site/transect location/trap location) to account for the non-independent spatial structure of the study design and used a Poisson error distribution. If the data were too sparse to fit three-way interactions of transect, habitat, and time, we fitted two-way interactions of transect and habitat as main fixed effects, and (1|site/transect location) as random effects. Note that “transect location” is a four-level factor referring to the spatial placement of each transect nested within a site, while “trap location” is a five-level factor referring to trap placement at five possible locations along a transect. We ran Wald tests and pairwise post hoc Tukey–Kramer tests to identify the statistical significance of fixed effects, and between-treatment response differences, respectively. We also

checked model fit by examining residual and fitted plots and checked for overdispersion by dividing the Pearson goodness-of-fit statistic by the residual degrees of freedom and ensuring values were below one (McCullagh and Nelder 1989). Data were analyzed using R 3.2.0 (R Development Core Team 2015), with the “lme4” (Bates et al. 2015), “car” (Fox et al. 2013), and “multcomp” (Hothorn et al. 2008) R packages for the GLMM analyses.

It is important to note that the split-plot design of our study meant that we were primarily interested in testing for a significant interaction of “transect” and “habitat” (definitions given in the previous section) to provide meaningful information on the effect of the specific farmland uses. In addition, we also were interested in testing for a “habitat” effect because it provides useful information on broader land-use differences between human-modified farmlands and natural remnant patches (see Fig. 1).

RESULTS

We collected a total of 4065 individual beetles, which comprised 280 species from 35 families (107 herbivore species, 100 predator species, and 73 detritivore species). The most abundant families were Anthicidae (10 species, $n = 1213$), Carabidae (48 species, $n = 757$), Staphylinidae (34 species, $n = 541$), Curculionidae (30 species, $n = 471$), and Tenebrionidae (25 species, $n = 383$; Appendix S1: Table S1).

There was adequate statistical power in the data for analyzing the responses of five families (Carabidae, Staphylinidae, Curculionidae, Anthicidae, and Tenebrionidae). Remnant patch size was discarded from the final models because the variation it explained was not significant (Appendix S1: Table S2).

Species composition

Overall beetle species composition was always significantly different between remnant patches and all farmland matrix types ($P < 0.03$) during spring, and between remnants and woody debris during summer ($P = 0.03$, Table 1). These compositional differences are further demonstrated with 92 species exclusively caught in remnant patches (e.g., *Cubicorhynchus* sp. #262 and *Georissus* sp.), 96 species in the farmland matrix (e.g., *Csiro* sp.), and

Table 1. Pairwise comparisons of beetle species composition dissimilarity (Bray–Curtis) between different land uses, based on permutational multivariate analysis of variance.

Habitat pairs	Spring			Summer		
	<i>F</i>	<i>R</i> ²	<i>P</i> adjusted	<i>F</i>	<i>R</i> ²	<i>P</i> adjusted
Patch vs. crop	3.07	0.06	0.015	2.31	0.05	0.090
Patch vs. plantings	1.98	0.04	0.030	1.01	0.02	1
Patch vs. fallow	2.02	0.04	0.030	1.92	0.04	0.120
Patch vs. woody debris	2.69	0.05	0.015	2.99	0.06	0.030
Crop vs. plantings	1.32	0.06	1	1.82	0.08	0.360
Crop vs. fallow	1.50	0.07	0.945	1.66	0.08	0.900
Crop vs. woody debris	0.34	0.02	1	1.14	0.06	1
Plantings vs. fallow	1.34	0.07	1	0.74	0.04	1
Plantings vs. woody debris	1.27	0.06	1	1.73	0.08	0.765
Fallow vs. woody debris	1.09	0.05	1	1.37	0.07	1

Note: Significant values ($P < 0.05$) are shown in bold.

92 species occurring in both habitats (e.g., *Omonadus hesperi* and *Gnathaphanus multipunctatus*, Fig. 2a). Within the farmland matrix, 14% of species occurred only in plantings, 14% only in fallows, 26% only in crops, and a large number of species (>26%) were shared between these different farmland uses (Fig. 2b). After woody debris was applied (only during summer), 36% of species were shared between the crop and woody debris (e.g., *O. hesperi*), while 27% occurred only in the woody debris (e.g., *Aridius* sp. #177 and *Longitarsus* sp. #272, Fig. 2c).

Species richness

Regardless of specific farmland matrix type, overall beetle species richness was significantly higher in the combined farmland matrix than in remnant patches on a per-trap basis (i.e., significant “habitat” effect, Table 2, Fig. 3a), with nearly twice as many species found in farmlands than in remnant patches. Species richness of predators, detritivores, and herbivores were not influenced by land use (i.e., effects of “habitat” and “habitat” × “transect” were non-significant, Table 2). Patterns of significantly higher species in the farmland matrix than in remnant patches were also exhibited by Carabidae, Staphylinidae, Anthicidae, and Tenebrionidae families (Appendix S1: Fig. S3a, b, e, h).

Between spring and summer, overall beetle species richness significantly decreased in all habitats (Fig. 3b).

Abundance

Responses for the abundance of a majority of beetle groups depended on interactive effects

“transect” × “habitat” × “time”, or “transect” × “habitat”. Specifically, we found significant three-way interactive effects of “habitat,” “transect,” and “time” on the abundance of overall beetles, predators, herbivores, as well as predatory Carabidae and herbivorous Curculionidae families. We found significant two-way interactive effects of “habitat” and “transect” on the abundance of detritivores, and the detritivorous Tenebrionidae family (Table 2).

Between spring and summer, predator abundance increased significantly in the fallow and planting matrix (Fig. 4a), while Carabidae abundance significantly decreased in the crop matrix and increased significantly in the fallow matrix (Fig. 5a).

Between spring and summer, herbivore abundance showed no significant differences among all farmland matrix types. Herbivore abundance decreased significantly between spring and summer in remnant patches adjacent to the crop, fallow, and plantings, but not in remnant patches adjacent to the woody debris (Fig. 4b). Curculionidae abundance was highest in fallow fields during spring and decreased significantly in the fallow and woody debris between spring and summer (Fig. 5b).

Detritivore abundance was significantly higher in the woody debris than in the adjacent remnant patch, but not significantly different when comparing crops and plantings with adjacent remnant patches. Detritivore abundance was significantly lower in the fallow than in the adjacent remnant patch (Fig. 4c). Tenebrionidae abundance was significantly higher in the fallow, planting, and

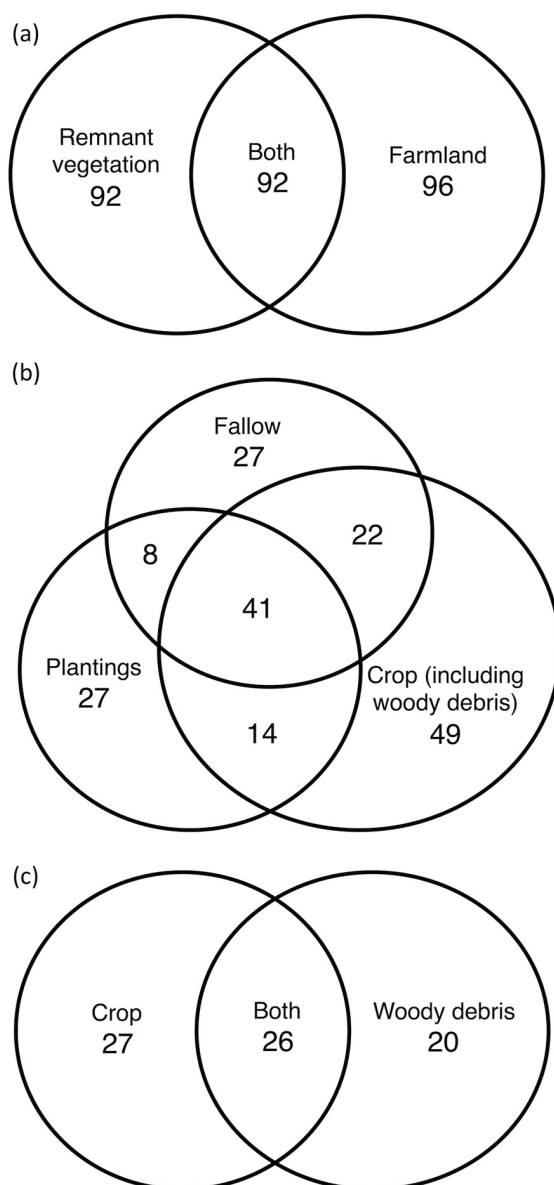


Fig. 2. Venn diagrams showing number of unique and shared species of beetles found (a) in remnant vegetation patch and farmland matrix habitats during both seasons; (b) in plantings, fallow, and crop fields (crop includes fields applied with fine woody debris) during both seasons; and (c) in crop and woody debris during summer only.

woody debris matrix than in the adjacent remnant patch, but differences between the crop matrix and adjacent remnant patch were non-significant (Fig. 5c).

DISCUSSION

We examined seasonal differences in beetle assemblages between woodland remnants and four contrasting farmland uses in a highly modified agricultural landscape. There were three key findings in our study: (1) Beetle species richness (per trap) was significantly lower in remnants than in all farmland uses combined; (2) beetle composition was significantly different between remnants and farmlands, with a third of species found in both habitats; and (3) abundance responses were often trophic group- or family-specific and influenced by interactions between land use and season. Our findings highlight the importance of maintaining a mosaic of land uses with both spatial and temporal heterogeneity to support beetle biodiversity in agricultural landscapes.

Higher species richness in farmlands than in woodland remnants

We found significantly higher beetle species richness in all farmland uses combined than in woodland remnants, on a per-trap basis. This pattern was underpinned by four abundant families of Carabidae, Staphylinidae, Anthicidae, and Tenebrionidae. This result was unexpected for this landscape because many arthropod taxa are associated with high levels of habitat complexity found in perennial native vegetation (i.e., habitat complexity hypothesis; Lassau et al. 2005, Attwood et al. 2008, Joern and Laws 2013) and are therefore adversely affected by habitat simplification and disturbance from intensive cropping and grazing land uses (Duelli et al. 1999, Hendrickx et al. 2007, Newbold et al. 2015). In addition, the majority of beetles sampled (>90%) were likely to be native species (K. Pullen, *personal communication*) and might be assumed to be associated with natural habitats. We also did not find abundance in farmlands represented by a few species, which is a pattern typically associated with higher productivity in agroecosystems (Ponce et al. 2011). Although our study did not directly examine specific mechanisms, we suggest three plausible interlinked reasons for why we found higher species richness in farmlands than in remnants.

The first possible reason for greater beetle richness in farmlands than in remnants is high nutrient inputs in farmlands, through fertilizer

Table 2. Summary of final generalized linear mixed models for species richness and abundance of (a) overall beetle assemblage, (b) trophic groups, and (c) common families, as predicted by transect type, habitat, time, or region.

Species richness				Abundance			
Model terms	χ^2	df	Pr ($>\chi^2$)	Model terms	χ^2	df	Pr ($>\chi^2$)
(a) All beetles							
Transect	3.11	3	0.375	Transect	4.44	3	0.218
Habitat	87.25	1	<0.001	Habitat	69.05	1	<0.001
Time	20.17	1	<0.001	Time	21.47	1	<0.001
Region	32.55	2	<0.001	Region	16.19	2	<0.001
Transect \times habitat	2.37	3	0.500	Transect \times habitat	20.91	3	<0.001
Transect \times time	3.99	3	0.263	Transect \times time	105.94	3	<0.001
Habitat \times time	2.65	1	0.104	Habitat \times time	0.15	1	0.703
Transect \times habitat \times time	6.07	3	0.108	Transect \times habitat \times time	21.70	3	<0.001
(b) Predators							
Transect	0.81	3	0.847	Transect	2.18	3	0.537
Habitat	2.27	1	0.132	Habitat	7.59	1	0.006
Time	1.34	1	0.247	Time	0.41	1	0.523
Region	6.48	2	0.039	Region	6.90	2	0.032
Transect \times habitat	0.08	3	0.994	Transect \times habitat	8.17	3	0.043
Transect \times time	1.69	3	0.640	Transect \times time	13.91	3	0.003
Habitat \times time	1.74	1	0.188	Habitat \times time	33.15	1	<0.001
Transect \times habitat \times time	1.54	3	0.674	Transect \times habitat \times time	19.33	3	<0.001
Detritivores							
Transect	1.63	3	0.652	Transect	1.29	3	0.733
Habitat	0.00	1	0.978	Habitat	0.71	1	0.400
Time	0.19	1	0.662	Time	49.43	1	<0.001
Region	5.24	2	0.073	Region	11.43	2	0.003
Transect \times habitat	1.61	3	0.656	Transect \times habitat	34.90	3	<0.001
Transect \times time	1.22	3	0.749	Transect \times time	12.29	3	0.006
Habitat \times time	0.05	1	0.818	Habitat \times time	17.13	1	<0.001
Transect \times habitat \times time	0.10	3	0.992	Transect \times habitat \times time	6.53	3	0.088
Herbivores							
Transect	5.51	3	0.138	Transect	4.11	3	0.250
Habitat	0.07	1	0.798	Habitat	0.27	1	0.602
Time	3.22	1	0.073	Time	81.88	1	<0.001
Region	3.34	2	0.188	Region	7.94	2	0.019
Transect \times habitat	2.69	3	0.442	Transect \times habitat	8.52	3	0.036
Transect \times time	1.99	3	0.574	Transect \times time	31.85	3	<0.001
Habitat \times time	0.03	1	0.863	Habitat \times time	12.20	1	<0.001
Transect \times habitat \times time	3.04	3	0.386	Transect \times habitat \times time	13.09	3	0.004
(c) Carabidae†							
Transect	6.45	3	0.092	Transect	10.28	3	0.016
Habitat	32.89	1	<0.001	Habitat	229.89	1	<0.001
Region	11.71	2	0.003	Time	63.96	1	<0.001
Transect \times habitat	4.52	3	0.211	Region	3.35	2	0.187
				Transect \times habitat	9.03	3	0.029
				Transect \times time	105.14	3	<0.001
				Habitat \times time	0.03	1	0.859
				Transect \times habitat \times time	18.04	3	<0.001
Staphylinidae†							
Transect	2.65	3	0.449	Transect	3.49	3	0.322
Habitat	14.48	1	<0.001	Habitat	22.56	1	<0.001
Region	5.54	2	0.063	Time	93.75	1	<0.001
Transect \times habitat	2.63	3	0.453	Region	7.41	2	0.025

(Table 2. *Continued*)

Model terms	Species richness			Abundance			
	χ^2	df	Pr ($>\chi^2$)	Model terms	χ^2	df	Pr ($>\chi^2$)
				Transect × habitat	4.33	3	0.228
				Transect × time	11.04	3	0.012
				Habitat × time	0.62	1	0.432
				Transect × habitat × time	3.97	3	0.264
Curculionidae†				Transect	10.78	3	0.013
Transect	7.59	3	0.055	Habitat	0.01	1	0.942
Habitat	0.53	1	0.466	Time	9.52	1	0.002
Region	2.85	2	0.240	Region	4.09	2	0.129
Transect × habitat	11.92	3	0.008	Transect × habitat	17.95	3	<0.001
				Transect × time	33.84	3	<0.001
				Habitat × time	0.01	1	0.912
				Transect × habitat × time	18.43	3	<0.001
Anthicidae				Transect	10.80	3	0.013
Transect	7.57	3	0.056	Habitat	59.60	1	<0.001
Habitat	32.19	1	<0.001	Time	148.35	1	<0.001
Time	1.61	1	0.204	Region	23.80	2	<0.001
Region	13.97	2	0.001	Transect × habitat	4.34	3	0.227
Transect × habitat	0.71	3	0.871	Transect × time	43.40	3	<0.001
Transect × time	1.51	3	0.679	Habitat × time	0.20	1	0.655
Habitat × time	0.23	1	0.635	Transect × habitat × time	6.20	3	0.102
Transect × habitat × time	0.70	3	0.872				
Tenebrionidae†				Transect	3.33	3	0.343
Transect	0.35	3	0.950	Habitat	94.63	1	<0.001
Habitat	9.15	1	0.002	Region	1.23	2	0.542
Region	1.89	2	0.389	Transect × habitat	18.60	3	<0.001
Transect × habitat	4.08	3	0.253				

Note: Significant values ($P < 0.05$) are shown in bold.

† Two-way interactions were fitted for species richness of families where data were too sparse to fit a three-way interaction.

use. This may have led to increased weed cover, which is a food resource for detritivores, as well as increased prey for insect predators (Abensperg-Traun et al. 1996). Other studies have, for example, found positive and unimodal relationships between productivity and species richness for arthropods and other taxonomic groups (Abensperg-Traun et al. 1996, Mittelbach et al. 2001).

Second, the dominance of extensive areas of farmlands (i.e., area effect) in the landscape may explain the high species richness in farmlands observed in our study (Fahrig 2003, Norton and Reid 2013). Increased resources and ecological niches in continuous farmland habitats may have led to more diverse assemblages, although species richness–area relationships in farmlands are strongly contingent on management practices in those areas (Norton and Reid 2013).

Lastly, the nature of predominant farming practices in the study region may have contributed to the persistence of many species in farmlands. Conservation tillage practices (i.e., reduced tillage and increased stubble retention, Llewellyn et al. 2012) have been adopted widely in Australian cropping systems over the past two decades, including in our study sites. The primary aim of these practices is to minimize soil loss (Holland 2004), but they may have had indirect conservation benefits for beetle assemblages. It is therefore possible that the biodiversity benefits from conservation tillage in Australia are comparable to “extensively managed” agroecosystems in Europe (Bennett et al. 2006, Kleijn et al. 2011) and Japan (Uchida and Ushimaru 2014). Several studies have found a large proportion of species adapted to early successional habitats associated with extensive farming practices (Duelli and Obrist

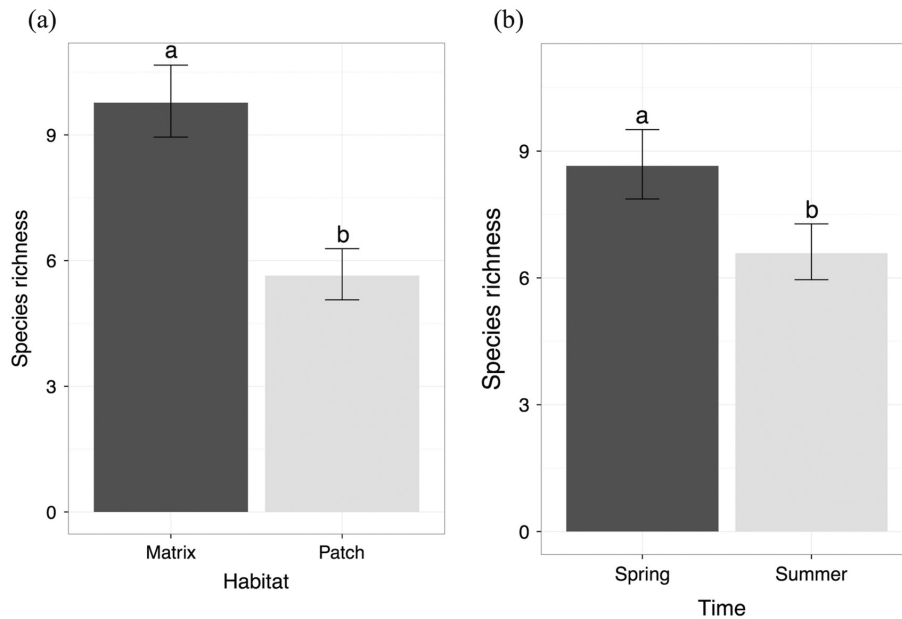


Fig. 3. Predicted mean species richness (per trap) of overall beetles by habitat (a) and time (b) (*P* values in Table 1). Patch refers to remnant vegetation, while matrix refers to four farmland uses combined (crop, fallow, planting, and woody debris). The 95% confidence intervals around predictions are shown. Different letters indicate significantly different results (Tukey–Kramer test).

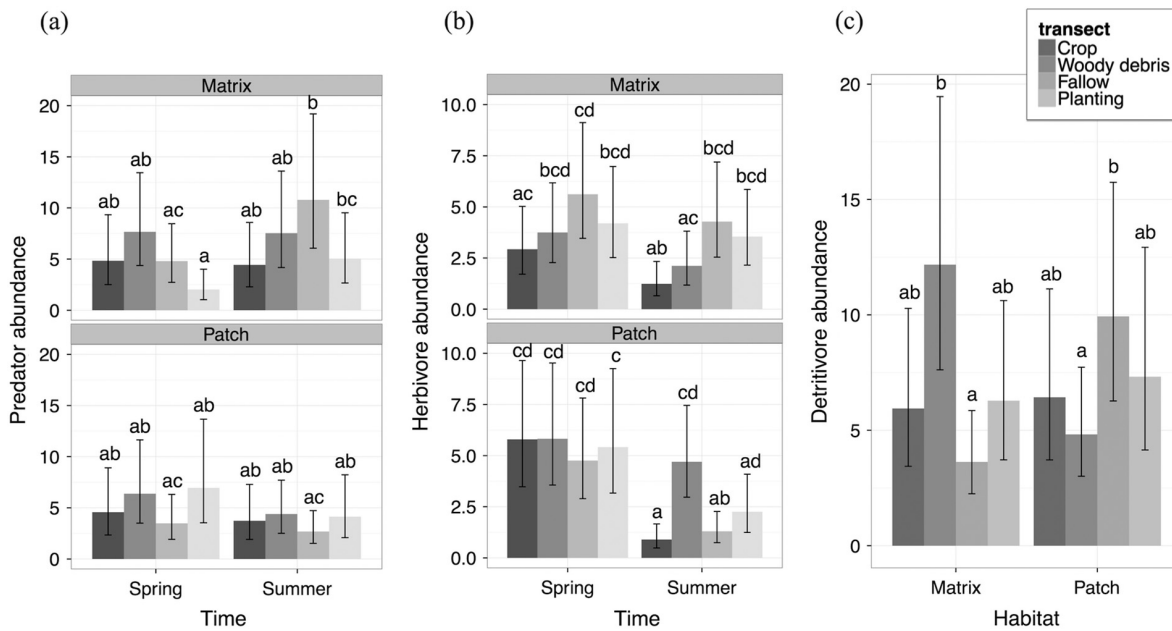


Fig. 4. Predicted mean abundance (per trap) of predators (a) and herbivores (b) to the interaction of “habitat,” “transect,” and “time,” and of detritivores (c) to the interaction of “habitat” and “transect” (*P* values in Table 1). Patch refers to remnant vegetation, while matrix refers to four farmland uses adjoining the remnant (crop, woody debris, fallow, and planting). The 95% confidence intervals around predictions are shown. Different letters indicate significantly different results (Tukey–Kramer test).

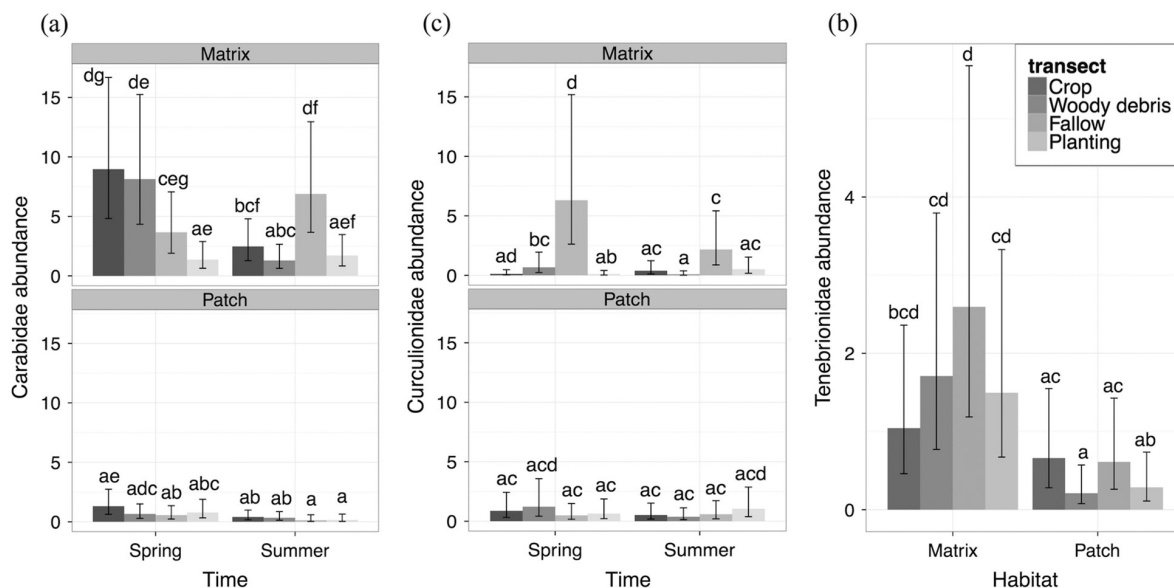


Fig. 5. Predicted mean abundance (per trap) of Carabidae (a) and Curculionidae (b) to the interaction of “habitat,” “transect,” and “time,” and of Tenebrionidae (c) to the interaction of “habitat” and “transect” (*P* values in Table 1). Patch refers to remnant vegetation, while matrix refers to four farmland uses adjoining the remnant (crop, woody debris, fallow, and planting). The 95% confidence intervals around predictions are shown. Different letters indicate significantly different results (Tukey–Kramer test).

2003, Bennett et al. 2006, Sutcliffe et al. 2015), which are characterized by moderate levels of disturbance and high levels of within-field spatial heterogeneity (Bennett et al. 2006, Kleijn et al. 2011, Uchida and Ushimaru 2014).

More studies are needed to determine whether further intensification of agricultural practices, such as management changes from conservation tillage to conventional tillage techniques, would result in declines in arthropod diversity. Long-term studies indicate limited adaptability of arthropod to high-intensity and high-frequency disturbance of soil (Stinner and House 1990, Lovei and Sunderland 1996). In Europe, intensification of farming practices in the 20th century, through increased mechanization, altered disturbance regimes, and the removal of remnant vegetation, has led to drastic reductions in arthropod biodiversity that previously inhabited extensive farming systems (Duelli and Obrist 2003, Tscharrntke et al. 2005, Sutcliffe et al. 2015).

Differences in species composition between land uses

A high proportion of beetle species were captured in a variety of farmland uses outside of

woodland remnants and appear to respond to spatial and temporal changes in resources or habitat quality associated with farmland heterogeneity. These results suggest that between-habitat heterogeneity (Benton et al. 2003, Vasseur et al. 2013)—particularly at the interface between farmland and remnants, and between different farmland uses—may be an important driver of assemblage patterns in mixed-farming landscapes.

We identified significant differences in species composition between farmlands and remnants (Table 2). Mechanisms underpinning beetle compositional differences could be explained by agricultural land-use changes, which have significantly modified native vegetation through introduction of exotic crops and pastures with agricultural intensification (Attwood et al. 2008, Newbold et al. 2015), and are consistent with the visualization in our principal components analysis showing contrasts in vegetation structure between remnants and farmlands (Appendix S1: Fig. S1). These compositional differences also likely reflect habitat specialization of some species, particularly open-habitat specialists in farmlands and woodland specialists in remnants (Thiele 1977). It should also be noted that the number of species unique to remnants was

comparatively higher than those unique to any single farmland uses, which suggests a potentially high number of specialists associated with remnant woodlands.

Spatio-temporal fluctuations of different beetle groups

Beetle species richness declined significantly in all land uses between spring and summer, a result that was consistent with predictions of stronger declines in species richness and abundance in more intensively managed land uses between spring and summer (Hill 1993, Sackmann and Flores 2009). Significantly lower species richness during late summer may be explained by life cycle dynamics (Sackmann and Flores 2009), with many species being less active or aestivating underground in adult or larvae form during hot summer conditions (Lovei and Sunderland 1996).

We observed interactive effects of land use and season on the abundance of all trophic groups and some families (i.e., Carabidae and Curculionidae), which supports previous work on the spatio-temporal dynamics of arthropods assemblages in agricultural landscapes (Benton et al. 2003, Vasseur et al. 2013). Different taxa have different habitat and resource requirements, which also change over time (Benton et al. 2003). We did not, however, observe stronger declines in abundance in more intensively managed land uses (Sackmann and Flores 2009). Our findings clearly show taxon-specific seasonal changes in habitat or food resources that were associated with specific farmland uses. Here, we discuss spatio-temporal abundance patterns exhibited by each beetle trophic group, and a representative family group, found in our study landscape.

Predators.—We found a significant increase in predator abundance in plantings and fallow fields between spring and summer, consistent with predictions of similar land uses having sufficient perennial elements as refuge during adverse summer conditions (Vasseur et al. 2013). For example, the abundance of Carabidae, which comprise the majority of predators in our study, likely followed peaks in resources between wheat crops during spring, and fallow fields during summer after crop harvest. This finding is consistent with studies in Europe and the United States, which identified wheat crops and weedy pastures as favorable

habitat and a source of weed seed or prey for polyphagous spring-breeding Carabidae (Lovei and Sunderland 1996, Kromp 1999, Woodcock et al. 2010). An absence of an increase in Carabidae abundance in plantings during summer, however, was unexpected because fallow fields and recent plantings have broadly similar ground layer complexity (K. Ng, *personal observation*). We suggest that other factors associated with plantings (e.g., predation on flightless ground beetles) might explain the inconsistent responses in fallow and plantings for Carabidae compared to predators more generally.

Detritivores.—We found that detritivores had similar or higher abundance in farmlands compared to remnant vegetation. Detritivore abundance was generally stable in farmlands and remnant vegetation, and significantly improved in woody debris when comparing with adjacent remnants. However, there was a possible negative effect of fallowing on detritivores. This may be linked to common management practices of fallow fields in this region, particularly grazing by livestock (Barton et al. 2011) or herbicide use (Baudry and Papyz 2001), which are associated with reduced beetle diversity. We also found that abundance of detritivorous Tenebrionidae could be augmented in farmlands by planting native vegetation, fallowing or applying woody debris on crop fields. Our results suggest that non-crop farmland uses may provide population sources of Tenebrionidae, in contrast with woodland remnants which have low numbers of Tenebrionidae. Some members of the Tenebrionidae family, such as *Adelium brevicorne*, *Isopteron* spp., and *Pterohelaeus* spp., are native pests of crop seedlings at larval stages (Gu et al. 2007, Micic et al. 2008). We suggest that retaining woodland remnants near crop fields may help reduce overall deleterious impacts of potential Tenebrionidae pests, although more work is needed to confirm this.

Herbivores.—Herbivore abundance, in general, did not differ significantly between remnants and farmlands during both spring and summer, which suggest that these contrasting land uses provide suitable host plant resources for different herbivore assemblages. Interestingly, there was a decrease in herbivore abundance in remnants between spring and summer, except for remnants adjacent to the woody debris treatment. We suggest that woody debris may mitigate

temporal decline of some herbivore species in remnants by increasing connectivity in crop fields, therefore facilitating colonization into remnants. Additional data would be needed to determine whether this pattern is related to observed declines in Curculionidae abundance in woody debris between spring and summer.

We also found higher abundance of herbivorous Curculionidae in fallow fields than other land uses during spring, which suggests that fallow fields provide optimal levels of spring-flowering weed or host plant resources for this family (Hangay and Zborowski 2010). This result is consistent previous studies which found high abundance of specialist Curculionidae species in grazed pastures (Batáry et al. 2007, Steiner et al. 2016), with the probability of occurrence for certain species increasing with grassland cover (Batáry et al. 2007). More research is needed to identify plant-species associations that may be driving high Curculionidae abundance when fallowing farmlands.

CONCLUSION AND MANAGEMENT IMPLICATIONS

Our findings contribute to growing evidence that effective conservation of arthropod diversity needs to consider entire landscape mosaics (Benton et al. 2003, Tscharrntke et al. 2005, Bennett et al. 2006, Vasseur et al. 2013) as well as maintain farmland heterogeneity with a mix of low-intensity land uses (Bennett et al. 2006, Kleijn et al. 2011, Uchida and Ushimaru 2014), such as conservation tillage, crop–fallow rotation, and restoration plantings. Complex taxon-specific abundance responses to interactions of land use and/or season indicate that no one single land use had optimal beetle diversity; rather, a diverse mix of farmland uses, which also consider the spatio-temporal heterogeneity of farm fields, is needed to conserve different beetle groups.

How farm fields are managed over time can influence patch dynamics and reduce extinction risks by providing complementary habitats or temporary connectivity for fragmented populations (Bennett et al. 2006, Driscoll et al. 2013, Vasseur et al. 2013). We demonstrated that applying fine woody debris to crop fields is a novel way of providing seasonal refuge for

detritivorous beetles (e.g., Tenebrionidae) and improving connectivity for herbivores (e.g., Curculionidae) without taking land out of food production. Fine woody debris may provide additional benefits, such as improving soil condition or providing resources for other litter-dependent fauna (Manning et al. 2013).

High overall species richness and abundance in farmlands suggest that farm fields can potentially be managed for both biodiversity conservation and agricultural production (Tscharrntke et al. 2005, Attwood et al. 2008, Bailey et al. 2010, Pywell et al. 2015). However, further intensification of agricultural practices in farmlands, such as increased monocropping, tillage, or agrochemical inputs, may undermine the high level of beetle biodiversity in this region (Tscharrntke et al. 2005, Cunningham et al. 2013, Sutcliffe et al. 2015).

Distinct assemblage composition in remnant vegetation patches indicates that farmland on its own is insufficient for conserving all beetle species. Retaining remnant vegetation is still critical for providing stable habitat and species persistence, especially for many species that depend on native vegetation (Bailey et al. 2010, Driscoll et al. 2013), are unable to survive agricultural disturbance, or use natural habitats at certain life stages (Thies et al. 2011, Driscoll et al. 2013).

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DATA AVAILABILITY

Data are available at CSIRO Data Portal: <https://doi.org/10.4225/08/5a00f9ab9a8d1>

SUPPORTING INFORMATION

Additional Supporting Information may be found online at: <http://onlinelibrary.wiley.com/doi/10.1002/ecs2.2042/full>