

**Exploring co-occurrence of closely-related guild members in a fragmented landscape  
subject to rapid transformation**

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1 **ABSTRACT**

2 Quantifying biotic responses to landscape transformation is a major research focus. Most past  
3 studies have explored co-occurrence of entire communities of a given group (e.g. birds)  
4 within largely intact ecosystems or over a limited time-frame. By contrast, here we use data  
5 from a 15-year experimental study, to explore intra-guild co-occurrence of six closely-related  
6 and functionally-similar sets of birds within 55 woodland fragments. Areas surrounding these  
7 remnants are undergoing transformation from grazed paddocks to *Pinus radiata* plantations,  
8 leading to a novel assemblage of forest and woodland birds. We sought to determine if the  
9 occurrence of a given species in a guild influenced the occurrence of other closely-related  
10 species in that guild, and through this relationship whether there was evidence of co-  
11 occurrence between species.

12 After controlling for environmental and habitat variables which can affect species occurrence  
13 like time since commencement of landscape transformation, patch size and vegetation type,  
14 we found the occurrence of a given species was influenced by the occurrence of a closely-  
15 related species in the same guild. Co-occurrence varied among bird guilds and included: (1)  
16 positive co-occurrence in which occurrence of one species within fragments positively  
17 affected the occurrence of another closely-related guild member (e.g. Eastern and Crimson  
18 Rosellas); and (2) negative co-occurrence in which the occurrence of one species was  
19 negatively associated with the occurrence of another within the same guild (e.g. Willie  
20 Wagtail and Grey Fantail).

21 We also identified interactions between patch size and species recording frequency within  
22 members of two guilds. For example, modelling of conditional recording frequency revealed  
23 the Eastern Rosella increased with increasing recordings of the Crimson Rosella in large  
24 patches, but decreased with increasing recordings of the Crimson Rosella in small patches.

25 Our results provide empirical evidence of co-occurrence among guild members and  
26 underscore the complexity of biotic responses to landscape transformation.  
27 **Keywords:** Co-occurrence; birds; landscape change; plantation; woodland fragments.

## 28 INTRODUCTION

29 The occurrence of species in human-modified landscapes can be influenced by an array  
30 of factors (Zuidema et al. 1996), many of which can be intimately inter-related (Lindenmayer  
31 and Fischer 2006). These include habitat loss (Fahrig 2003), habitat fragmentation (Collinge  
32 2009), temporal changes in habitat suitability (Felton et al. 2003), the size of remaining  
33 patches of habitat (Hanski 1994), edge effects (Ries et al. 2004), the condition of and changes  
34 in the matrix surrounding habitat patches (Driscoll et al. 2013), and altered key ecosystem  
35 processes (Galetti et al. 2013).

36 Species occurrence in rapidly transforming landscapes also may be influenced by the  
37 presence and abundance of other species (Maron and Kennedy 2007, Ovaskainen and  
38 Soininen 2011, Godsoe and Harmon 2012, Mac Nally et al. 2012). This may arise because of  
39 increased aggressive interactions between species (Mac Nally et al. 2012), increased levels of  
40 parasitism (Smith et al. 2003), and disrupted mutualisms (Bascompte and Jordano 2007).  
41 However, most studies of co-occurrence focus on entire communities of a given group of  
42 organisms (e.g. birds or mammals), have not taken place in landscapes undergoing rapid  
43 transformation, and/or are conducted either at one point in time or over a limited time-frame  
44 (2-3 years) (but see Heske et al. 1994, Sebastian-Gonzalez et al. 2012). Here we take a  
45 different approach by exploring the tendency of two species to occur together (or apart), after  
46 controlling for other factors that affect the occurrence of the species. We do this using  
47 systematic bird surveys from a 15 year observational study in a rapidly transforming  
48 landscape to examine co-occurrence among sets of closely-related, morphologically-similar  
49 and functionally-similar bird species.

50 Our study area was characterized by 55 fragments of temperate eucalypt woodland  
51 around which the landscape was undergoing marked and ongoing transformation from grazed  
52 paddocks to stands of densely-spaced Radiata Pine (*Pinus radiata*) plantation trees. The

53 landscape is therefore a mosaic (sensu Bennett et al. 2006) of woodland patches embedded  
54 within an extensive exotic plantation. In Australia, temperate eucalypt woodland  
55 ecosystems are recognized as a distinctly different environment from native eucalypt forest  
56 ecosystems because of differences in: **(1)** dominant tree, understorey and ground cover plants,  
57 **(2)** vegetation structure (Specht and Specht 1999), and **(3)** animal biota (Lindenmayer et al.  
58 2010). In earlier work, we found that some forest-associated bird species have colonised the  
59 exotic Radiata Pine stands and then spilled over to also occupy adjacent woodland fragments  
60 (Lindenmayer et al. 2008). As a result, the woodland fragments support a novel assemblage  
61 (sensu Hobbs et al. 2006) of woodland bird and forest bird species that does not occur  
62 elsewhere in the region's forests or in the region's temperate woodlands (Lindenmayer et al.  
63 2008). For a number of bird guilds, the woodland species and the forest species are closely  
64 related and morphologically and functionally similar (see Appendix 1). Given this, the  
65 primary question which underpinned our investigation was:

66 *Is the occurrence of a given species in a guild influenced by the occurrence of another closely*  
67 *related species in that guild?*

68 We recognized three possible broad kinds of co-occurrence for this study. These were:  
69 **(1) Positive co-occurrence** in which the occurrence within woodland patches of one species in  
70 a given guild had a positive effect on the occurrence of another closely related species in that  
71 same guild. **(2) Negative co-occurrence** in which the presence of one species was negatively  
72 associated with the presence of another within the same guild. And **(3) An absence of positive**  
73 **or negative patterns of co-occurrence**. That is, where there was no evidence of significant  
74 positive or negative influence of one species on the occurrence of one or more members of  
75 the same guild.

76 Theory and empirical studies suggest the existence of positive co-occurrence in some  
77 ecosystems but negative co-occurrence in others (Ovaskainen et al. 2010, Andersen et al.

78 2013). Positive co-occurrence might occur because functionally similar and/or closely related  
79 taxa might be adapted to similar environments (Ovaskainen et al. 2010) or gain mutual  
80 benefits (like enhanced foraging opportunities in mixed-species feeding flocks) (Bell 1980,  
81 Sridhar et al. 2012). An additional reason for positive co-occurrence could be that a given  
82 species might choose habitat using information gleaned from other species present at a  
83 location (Smith and Hellman 2002, Sebastian-Gonzalez et al. 2012), particularly a species  
84 that is very similar to itself (Seppänen et al. 2007). Alternatively, two or more  
85 morphologically and/or functionally similar species may exhibit negative co-occurrence  
86 because of competition (e.g. see Alatalo et al. 1986, Lovette and Hochacka 2006, Steen et al.  
87 2014), particularly when resources are limited (Newton 1998). Given the potential  
88 manifestation of processes such as competition versus inter-specific attraction, we focused  
89 this study on six closely-related, morphologically similar, and functionally similar guilds of  
90 species to maximize the chances of detecting positive or negative co-occurrence (see Gaston  
91 1996).

92         We were acutely aware that two species within a guild may co-occur in an area for  
93 reasons other than the influence of one species on the other (e.g. by having similar habitat  
94 requirements (Ovaskainen et al. 2010)). Therefore, a novel feature of our study was to  
95 explore patterns of co-occurrence while accounting for habitat and environmental variables  
96 that we believe also may influence the occurrence of particular species. These included  
97 vegetation type and the time elapsed since the start of transformation of the landscape  
98 surrounding the remnant woodland patches. We also recognized that the potential for the  
99 occurrence of one species to affect the occurrence of another might be influenced by factors  
100 like patch size. This may arise because, for example, large patches may support more  
101 resources (see Zarette et al. 2000) or more heterogeneous than small patches (Lindenmayer  
102 and Fischer 2006). More resources and/or greater heterogeneity may make it possible for

103 closely related species to co-exist (Jeltsch et al. 1998) such as in large heterogeneous patches  
104 but not in smaller, more homogeneous patches. Therefore, a second question we addressed in  
105 this study was:

106 *If the presence of a given species is influenced by the occurrence of another closely-related*  
107 *species from the same guild, does that relationship change with patch size? That is, is there*  
108 *evidence of an interaction between the occurrence of a predictor species and patch size?*

## 109 **METHODS**

### 110 **Study area**

111 Our study region is called the Nanangroe area and it is 10-20 km south-east of the town  
112 of Jugiong in southern New South Wales, south-eastern Australia (Fig. 1). Nanangroe lies  
113 approximately 70 km north-west of Canberra (coordinates 34.9°–35.0°S and 148.4°–148.5°E,  
114 altitudinal range: 250-750 m asl) and is characterised by a temperate climate (hot summers  
115 and relatively cool winters). The original vegetation cover in the study area included several  
116 temperate woodland vegetation types. Over 80 per cent of the vegetation cover in the study  
117 area has been cleared in the past 150 years, primarily for domestic stock grazing (McKernan  
118 2010). Vegetation in these heavily cleared paddocks is dominated by exotic pasture grasses  
119 and isolated single trees that are the remnants of past woodland vegetation cover. Prolonged  
120 periods of livestock grazing in the region meant that understorey vegetation of *Acacia* spp.  
121 and other plants was largely absent from all areas of woodland in the study area. In several  
122 parts of the region, an emerging pine plantation now surrounds patches of remnant woodland  
123 that escaped earlier waves of land clearing. Further details of the study area can be found in  
124 Lindenmayer et al. (2008).

### 125 **The design of the Nanangroe study**

126 In 1998, prior to commencement of landscape transformation to a pine-dominated  
127 system, we randomly selected 55 of the 70 remnant woodland patches stratified by vegetation

128 class and woodland patch size. These woodland patches were exempt from clearing while  
129 plantation establishment in the surrounding areas was undertaken. The 55 remnant woodland  
130 patches varied in dominant vegetation type, including Yellow Box (*Eucalyptus melliodora*),  
131 Red Box (*E. polyanthemos*), White Box (*E. albens*), Blakely's Red Gum (*E. blakelyi*), Apple  
132 Box (*E. bridgesiana*) and Long-leaf Box (*E. goniocalyx*). In addition, there were patches  
133 dominated by Red Stringybark (*E. macrorhyncha*), Broad-leaved Peppermint (*E. dives*), and  
134 River Oak (*Allocasuarina cunninghamiana*).

135         The size of the patches in our study ranged from 0.5 to 28.8 ha (median = 1.5 ha, 32  
136 remnants < 2 ha) and they were typically located at least 500 m apart. Although our  
137 investigation included some larger remnant patches (e.g. eight > 5 ha), numerous large  
138 woodland patches were not available for study because of the extent of past land clearing for  
139 agriculture and domestic livestock grazing in this part of Australia (McKernan 2010).  
140 However, we note that over the past decade, in many patches, including the small patches,  
141 there is strong evidence of breeding by many of the 15 species of birds we have targeted in  
142 this study.

143         Between 1997 and 2011 grazing by domestic livestock continued throughout the pine  
144 plantation and the 55 patches of remnant woodland at Nanangroe. This was important to  
145 ensure that grazing and landscape context effects were not confounded.

#### 146 **Bird surveys**

147         To survey birds, we established a permanent 200 m long transect randomly within each  
148 of the 55 woodland patches. For each survey, two observers each visited three points at 0 m,  
149 100 m and 200 m along the transect on different days; typically 2-4 days apart. At each point,  
150 the observer recorded which species were detected in a 5 minute interval within the woodland  
151 patch and within 50m of the point. The above layout of point-interval count stations within a  
152 site was consistent across the study except in some of small patches where the third station



153 was placed closer than 100m to the second station. In the few instances where a portion of the  
154 station's 50m radius was not confined within the woodland patch, our field protocols  
155 restricted the recording of birds to only those within the patch (i.e. not in the adjacent pine  
156 plantation). We completed field surveys in early November of the following years- 1999,  
157 2001, 2003, 2005, 2007, 2009, and 2011. Early November is the peak breeding season in the  
158 study region, when summer migrants are present and birds have established territories and  
159 exhibit strong patterns of site fidelity (Lindenmayer et al. 2008). Detections were made  
160 between 5.30-9.30am and were not undertaken on days of poor weather (rain, high wind, fog  
161 or heavy cloud cover). The order in which particular sites were surveyed on any given day  
162 was varied so that if birds were sampled, for example, late in the morning on a site on one  
163 day, that site was surveyed early in a subsequent survey.

164 Bird observers with a minimum of ten years of experience completed surveys of forest  
165 and woodland birds. These observers varied in their ability to detect some groups of birds but  
166 Lindenmayer et al. (2009) showed that pooling counts of two or more observers at the same  
167 plot point could compensate for extra variability due to observer heterogeneity. The 15  
168 species we investigated in this study were readily recognizable taxa with distinctive calls.  
169 Moreover, we were familiar with them from many previous studies in a range of  
170 environments in south-eastern Australia (Lindenmayer et al. 2009). We therefore assumed  
171 that mis-identification was low for the species in the six guilds included in this study.

## 172 **Bird guilds**

173 We selected six groups of birds for study using three key criteria. First, the members of  
174 a group were closely related – almost always congeneric, always from the same family, and  
175 always from the same dietary guild (see Appendix 1). Birds in the same group also were  
176 similar in body size. Second, the six groups represented different foraging guilds, diets,  
177 breeding strategies and other life history attributes. Therefore, our analyses enabled us to

178 determine if patterns of co-occurrence were consistent across groupings. A third criterion for  
179 selection was that at least one member of a given set of birds was a woodland or open-  
180 country species, whereas one or more of the others was a forest-associated species This  
181 enabled us to explore whether there was avoidance by woodland-associated taxa of forest  
182 birds in the respective groups of birds as the pine plantation stands surrounding the woodland  
183 patches matured.

#### 184 **Bird data used in statistical analyses**

185 We obtained bird species presence/absence data by pooling the six point-interval counts per  
186 site for each survey year. We also examined a recording frequency index which was the  
187 proportion of plots (out of a maximum of six in any given survey year) at which a given  
188 species was recorded.

189 We made comparisons of bird taxa only within groups of similar species so the  
190 interpretation of the recording frequency was the same for each species in a given group.  
191 Finally, we used the same field methodology through time so the comparisons within groups  
192 of similar species were meaningful.

#### 193 **Statistical analyses**

194 We conducted our statistical analyses in a series of steps. First, we completed graphical  
195 analyses to examine patterns in species occurrence and our recording frequency in relation to  
196 plantation age, vegetation type and the presence and abundance of other guild member  
197 species. Second, we used a univariate two-part binomial model to examine the factors  
198 influencing the occurrence and conditional abundance of birds. Third, we fitted the Bayesian  
199 multivariate logistic regression model of O'Brien and Dunson (2004) to determine the  
200 consistency of results obtained from the univariate two-part models.

201 Welsh et al. (1996) used a two-part model to examine fauna abundance data. The first  
202 component of their model estimates the probability of a species being present using binary

203 logistic regression and the second component models abundance, conditional on the animal  
204 being present. The two-part model was developed using a zero-truncated Poisson or Negative  
205 Binomial distribution for the second (conditional) part of the model (Welsh et al. 1996). We  
206 have not measured abundance directly in this investigation, but adopt this general modelling  
207 framework and substitute a truncated Binomial distribution for the second part of the model  
208 to describe the recording frequency in lieu of a direct measure of abundance. We refer to the  
209 second part of the model as the (mean) conditional recording frequency. The `vglm()` function  
210 in the R VGAM package was used, with the second conditional part of the model fit using the  
211 `posbinomial` (zero-truncated binomial) distribution family ([see Appendix 2 and Yee 2012](#)).  
212 The two-part modelling approach allows for over-dispersion and both parts of the model  
213 allow for serial dependence by including the lagged value of the response as an explanatory  
214 variable. The lagged value is the response recorded in the previous survey two years before—  
215 lagged presence for the first part of the model and lagged recording frequency for the second  
216 part of the model. We used a lagged value of two years to account for the fact that our field  
217 surveys were conducted two years apart.

218 A number of factors can influence species response in a complex, fragmented  
219 agricultural environment. To more clearly identify the role that fellow-guild member species  
220 may play in species occurrence (and through this, co-occurrence), it was advantageous to  
221 include some key environmental covariates in the analysis. We fitted four types of two-part  
222 binomial models which differed in their complexity or number of covariates.

223 **(1) The base model** included the lagged value of the response, guild member predictor  
224 species, years since planting, and log patch size. The idea was that the potential explanatory  
225 variables for guild member species captured positive (or negative) co-occurrence with a  
226 positive coefficient (or a negative coefficient) and the years since planting captured change  
227 through time as the Radiata Pine plantation surrounding the woodland patches matured. With

228 respect to the guild member predictor species term in the model, species presence is used for  
229 the first (occurrence) part of the model and the recording frequency is used for the second,  
230 conditional recording frequency stage. For all models, the focus was on the occurrence part of  
231 the model but we also were interested to see if the conditional recording frequency revealed  
232 any patterns among the species.

233 **(2) The vegetation model** included the same terms as the base model but also included a four  
234 level factor for vegetation type which described the dominant vegetation community of the  
235 remnants. The possible values for vegetation community type were: **(i)** Scribbly Gum  
236 (*Eucalyptus rossii*)-stringybark (*E.macrorhyncha*)-silvertop ash (*E. sieberi*); **(ii)** Swamp Gum  
237 (*E. ovata*)-Snow Gum (*E. pauciflora*); **(iii)** Yellow Box (*E. melliodora*)-Blakely's Red Gum  
238 (*E. blakelyi*)-River Oak (*Casuarina cunninghamiana*); and **(iv)** a White Box (*E. albens*)  
239 dominated community with associated Yellow Box and Blakely's Red Gum. Vegetation type  
240 was coded to provide contrasts for each level against the baseline reference level Scribbly  
241 Gum-stringybark-silvertop ash.

242 **(3) The interaction model** included an interaction term between guild member predictor  
243 species (presence or conditional recording frequency) and log patch size in addition to the  
244 terms included in the base model. And,

245 **(4) A composite model** which included vegetation type and the interaction terms, as well as  
246 the terms included in the base model. We used the Bayesian (Schwarz) information criteria  
247 (BIC) to choose between the four types of models for each bird species response (Schwarz  
248 1978) (Appendix 4A).

249 For the univariate two-part models, we interpreted co-occurrence as significant  
250 coefficients in paired sets of regressions within a set of guild models. For example, within a  
251 guild, species B and C may have a positive effect on the probability of occurrence of the  
252 response, species A. In a second, corresponding model, the probability of occurrence for

253 species B may be positively influenced by species A but not species C (*viz*: an absence of co-  
254 occurrence). Here, we interpret this complementary result as evidence that species A and B  
255 co-occur. The models will have one less predictor species than there are guild members: for a  
256 guild with two members like the rosellas, there will be one predictor species and for the four-  
257 member thornbill guild there will be three predictor species in each model. We note that the  
258 second, conditional recording frequency part of the two-part model may also provide  
259 evidence of co-occurrence under this interpretation, where a matching or reciprocal  
260 relationship in regression coefficients is present.

261 The log patch size variable and vegetation type were of interest because occurrence and  
262 recording frequency may depend on the size and dominant vegetation of a patch. As is usual  
263 in transition models for longitudinal data, the lagged response variable was intended to  
264 capture serial dependence and was not of interest in its own right. We examined spatial  
265 dependence among sites for each response variable using variograms of the Pearson residuals  
266 from each of the logistic regression models.

267 In the Bayesian multivariate logistic regression analysis, we modelled the joint  
268 probability of occurrence of each species in a given guild. For each response component, we  
269 fitted the lagged species occurrence indicator (as defined above for the two-part model) for  
270 each species to capture the serial dependence, the years since planting of the surrounding  
271 Radiata Pine plantation, and the logarithm of the patch size. Here, the co-occurrence effects  
272 were obtained directly from the pairwise correlations between the different response  
273 components on the logistic scale (O'Brien and Dunson 2004). For other applications of the  
274 approach taken by O'Brien and Dunson (2004), see Ovaskainen et al. (2010) and [Sebastian-  
275 Gonzalez et al. \(2012\)](#). Bayesian *P*-values (bpv) were reported to provide a measure of the  
276 strength of evidence for individual terms in the model. All statistical analysis was done using  
277 R (R Development Core Team 2013).

## 278 RESULTS

279 We summarize our data on the number of woodland patches occupied by each of the 15  
280 species in the six guilds in each survey year in Table 1. These data highlight marked  
281 interspecific differences and temporal changes in the number of occupied patches. Our  
282 graphical approach revealed patterns in occurrence and recording frequency between species  
283 within a guild. For example, there was increasing recording frequency of the Brown  
284 Thornbill with increasing time since planting with some evidence that the rate or magnitude  
285 of increase may differ among vegetation types. The frequency of recording of the Yellow-  
286 rumped Thornbill appeared to decrease with increasing time since planting (Appendix 3).

287 The use of BIC to select from our four main groups of models resulted in the base  
288 model being preferred in all but a few cases (see Appendix 4A). The results of univariate  
289 two-part modelling provided evidence for positive co-occurrence in five pairs of species,  
290 negative co-occurrence in one pair of species, and an absence of (positive or negative) co-  
291 occurrence patterns in seven pairs of species. Positive co-occurrence occurred among  
292 members of three guilds – the rosellas, flycatchers, and thornbills. Conversely, we found  
293 evidence of negative co-occurrence among two members of the flycatcher guild. There was  
294 no discernible pattern of (positive or negative) co-occurrence for the two members of the  
295 treecreeper group, nor for the two members of the whistler group.

### 296 *Rosellas*

297 For the rosellas, the presence/absence component of the two-part model contained  
298 evidence of co-occurrence (Fig. 2), and the probability of occurrence of the Eastern Rosella  
299 decreased with years since planting ( $p=0.019$ , Fig. 2). For the Eastern Rosella, we identified  
300 an interaction between log patch size and recording frequency of the guild member species  
301 the Crimson Rosella ( $p<0.002$ ); as the recording frequency of the Crimson Rosella increased,

302 the conditional recording frequency of the Eastern Rosellas fell for small patches but  
303 increased in larger patches.

#### 304 *Flycatchers*

305 In the flycatcher group, the Willie Wagtail and the Restless Flycatcher exhibited  
306 positive patterns of co-occurrence while the Willie Wagtail and the Grey Fantail showed  
307 negative co-occurrence (Fig. 3).

#### 308 *Thornbills*

309 From the presence/absence component of the two-part models, we inferred patterns of  
310 co-occurrence among the Striated Thornbill and the Brown Thornbill, the Striated Thornbill  
311 and the Buff-rumped Thornbill, and the Buff-rumped Thornbill and the Yellow-rumped  
312 Thornbill (Fig. 4). There was a positive interaction between log patch size and the guild  
313 member predictor species – the Striated Thornbill ( $p < 0.005$ ) and the Yellow-rumped  
314 Thornbill ( $p = 0.03$ ). As the numbers of the Striated or Yellow-rumped Thornbill increased,  
315 the conditional recording frequency of the Buff-rumped Thornbill decreased in small patches  
316 but increased within large patches. There was no evidence of a reciprocal interaction for  
317 either the Striated or Yellow-rumped Thornbill. We note that both the Brown Thornbill and  
318 the Buff-rumped Thornbill, and the Striated Thornbill and the Yellow-rumped Thornbill  
319 exhibited no evidence of co-occurrence.

#### 320 *Honeyeaters*

321 Although the paired regression coefficients were not significant for the honeyeaters  
322 (indicating co-occurrence), we found weak evidence that the probability of occurrence of the  
323 White-plumed Honeyeater was reduced when the Yellow-faced Honeyeater was present  
324 ( $p = 0.064$ , Fig. 2) and that the conditional recording frequency of the Yellow-faced  
325 Honeyeater decreased as the conditional recording frequency of the White-plumed  
326 Honeyeater increased ( $p = 0.013$ ).

327 Results obtained from fitting Bayesian multivariate logistic regression models (as per  
328 O'Brien and Dunson 2004) were broadly consistent with those from the univariate two-part  
329 binomial models and are summarized in Figure 5 and supplementary material (Appendices  
330 4D and 4E)

331 We have focused the reporting of our results on evidence for co-occurrence. However,  
332 our modelling also indicated that the presence and/or conditional recording frequency of  
333 almost all species also was significantly influenced by factors such as time since planting, log  
334 patch size, vegetation type or the presence and/or conditional abundance of fellow-guild  
335 member species. The full results are listed in Appendices 3B and 3C. Examination of  
336 variograms of Pearson residuals revealed no substantial, systematic spatial dependence  
337 among the sites in our dataset.

## 338 **DISCUSSION**

339 Landscape change and habitat fragmentation are major land management issues  
340 worldwide and understanding their impacts on biota is critical for developing informed  
341 strategies for biodiversity conservation (Lindenmayer and Fischer 2006). Similarly,  
342 quantifying patterns of co-occurrence and understanding the factors that influence such  
343 patterns remains one of the most enduring and controversial issues in ecology (Sfenthourakis  
344 et al. 2005, Ovaskainen et al. 2010, Sridhar et al. 2012, Andersen et al. 2013, Steen et al.  
345 2014). Indeed, Dammhahn and Kappeler (2008, p. 473) noted that” *Understanding the co-*  
346 *occurrence of ecologically similar species remains a puzzling issue in ecology*”.

347 In this study, we have examined a topic that lies at the intersection of these two  
348 substantial arenas of ecological research by exploring co-occurrence among closely related  
349 (intra-guild) sets of bird species in a landscape subject to major and rapid change. Our  
350 primary question was: *Is the occurrence of a given species in a guild influenced by the*  
351 *occurrence of another closely related species in that guild?* The answer to this question was



352 yes for some of the members of several guilds– with the results of our landscape experiment  
353 indicating that, after controlling for key factors like the time elapsed since landscape  
354 transformation, there were effects of the presence or recording frequency of a given species  
355 on the occurrence of other, closely-related and functionally-similar species in the same guild.  
356 The existence of positive or negative co-occurrence varied among some sets of species and  
357 also between species within the same guild, including for those cases where three or more  
358 taxa were considered.

359 An interesting result of our investigation was an interaction between patch size and  
360 recording frequency for the members of two bird guilds; with the conditional recording  
361 frequency of one species in a group increasing with increasing frequency of another member  
362 of the guild in large patches, but decreasing with increasing frequency in small patches. Thus,  
363 the answer to our second question posed at the outset of this study – *Does the influence that*  
364 *one species has on the presence of another change with patch size?* – also was yes, albeit not  
365 for all species. We further discuss these and other findings in the remainder of this paper.

## 366 **Broad kinds of co-occurrence**

### 367 *Positive co-occurrence patterns*

368 A key aim of our study was to explore the types of co-occurrence among species in the  
369 same guild in a rapidly transforming landscape, while at the same time controlling for other  
370 factors which can influence species like time since landscape transformation commenced,  
371 patch size and vegetation type. This novel environmental and habitat filtering approach  
372 provided evidence of positive co-occurrence in three groups – the rosellas, flycatchers and  
373 thornbills. For example, the probability of occurrence of the Eastern Rosella increased  
374 significantly when the Crimson Rosella was present and the reverse (i.e. reciprocal) pattern  
375 also was present. That is, the probability of occurrence of both species was positively  
376 influenced by the presence of the other in the respective, paired models. Similarly, the

377 probability of occurrence of the Striated Thornbill increased significantly when the Brown  
378 Thornbill was present. The complementary relationship also occurred. That is, the probability  
379 of occurrence of the Brown Thornbill increased significantly when the Striated Thornbill was  
380 present. Similar complementary relationships to those for the Striated and Brown Thornbills  
381 were found for the Striated and Buff-rumped Thornbills, and the Buff-rumped and Yellow-  
382 rumped Thornbills.

383 Several factors may explain the positive co-occurrence patterns that we observed in this  
384 study. In the case of the thornbills, earlier studies of this group have shown that several  
385 species of thornbills often form mixed feeding flocks (Bell 1980). Sridhar et al. (2012)  
386 showed in an analysis of a global dataset that positive patterns co-occurrence can arise among  
387 bird species which share similar taxonomy, body size and foraging methods. This may occur  
388 because of co-operative benefits like the discovery of food resources or recognition of alarm  
389 calls – that result from being a member of such kinds of flocks, especially when there is a  
390 high level of similarity among species (Sridhar et al. 2012). Notably, other studies have found  
391 that among some species such as breeding waterbirds, there is heterospecific attraction  
392 leading to positive co-occurrence (Ovaskainen et al. 2010). Such patterns may arise because  
393 of calling behaviour that alerts another species to the presence of predators or the location of  
394 high quality habitat (Catchpole and Slater 1995, Smith and Hellman 2002). An additional or  
395 alternative explanation for positive co-occurrence could be that a given species might choose  
396 habitat using information gleaned from other species present at a location (Smith and  
397 Hellman 2002, Sebastian-Gonzalez et al. 2012). This is particularly true for a species that is  
398 very similar to itself (Seppänen et al. 2007) as occurs for many of the intra-guild members  
399 examined in this investigation. However, a novel feature of our study was to explore patterns  
400 of co-occurrence while accounting for habitat and environmental variables. This was  
401 important as two species within a guild may co-occur because of similar habitat requirements

402 (Ovaskainen et al. 2010) and thus we attempted to account for this prior to exploring the  
403 influence of one or more intra-guild members on another guild member. *Negative co-*  
404 *occurrence patterns*

405 We recorded negative co-occurrence among two species in the flycatcher guild. We  
406 found that after accounting for log patch size and years since planting, the probability of  
407 occurrence of the Grey Fantail declined when the Willie Wagtail was present. The reciprocal  
408 relationship also occurred in which (after accounting for other explanatory variables), the  
409 occurrence of the Willie Wagtail declined when the Grey Fantail was present. Both these  
410 species can be relatively aggressive and are known to attack other birds and this may explain  
411 the negative co-occurrence or avoidance patterns we observed in this study. Notably, such  
412 relationships were not observed with the third member of the flycatcher guild – the Restless  
413 Flycatcher, suggesting asymmetrical co-occurrence patterns among group members  
414 reinforced by the positive co-occurrence inferred from the model between the Willie Wagtail  
415 and Restless Flycatcher. The Willie Wagtail and Restless Flycatcher are typically woodland  
416 birds whereas the Grey Fantail is generally a forest-associated species. Indeed, the Willie  
417 Wagtail and the Restless Flycatcher are common co-inhabitants of remnants of temperate  
418 eucalypt woodland throughout large parts of inland eastern Australia. Conversely, it is  
419 possible that the development of the novel ecosystem in the Nanangroe area has led to an  
420 aggressive woodland species (the Willie Wagtail) avoiding an aggressive forest species (the  
421 Grey Fantail).

422 There was weak evidence that the probability of occurrence of the White-plumed  
423 Honeyeater was reduced when the Yellow-faced Honeyeater was present and that the  
424 conditional recording frequency of the Yellow-faced Honeyeater decreased as the conditional  
425 recording frequency of the White-plumed Honeyeater increased. These two species have  
426 similar habitat, nesting and dietary requirements and it is possible that the negative

427 relationships in conditional abundance and occurrence (Fig. 2) reflects competition between  
428 them. Notably, other authors have identified negative associations between the Yellow-faced  
429 and White-plumed Honeyeaters (Kinross and Nicol 2008), although unlike this investigation,  
430 they did not account for the effects of other important habitat-related factors such as  
431 vegetation type that can influence occurrence.

432 We suggest that because we have controlled for factors like vegetation type in our  
433 study, then the negative intra-guild co-occurrence patterns we observed are likely to be driven  
434 by competition and resource availability. Transformation of the Nanangroe landscape has  
435 altered the matrix around the remnants, and likely then altered the resources available to  
436 species within the patches (Driscoll et al. 2013). This leads to altered competitive dynamics  
437 among similar species, leading to “winners and losers” and in turn, changes in the bird  
438 community over time to produce the novel assemblages (sensu Hobbs et al. 2013) observed  
439 in the Nanangroe landscape (Lindenmayer et al. 2008).

#### 440 **Patch size effects**

441 Our analyses revealed a significant interaction between log patch size and conditional  
442 recording frequency for the Eastern Rosella and the Buff-rumped Thornbill with respective  
443 guild members. In both cases, the patterns of increasing conditional recording frequency that  
444 were accompanied by increasing recording frequency of guild member species in large  
445 patches of woodland surrounded by maturing pine stands were reversed in small patches. The  
446 reasons for these patch size effects remain unclear. However, at the outset of this study, we  
447 postulated (see Question 2) that interactions between patch size and occurrence and/or  
448 recording frequency patterns may occur because large patches may be more resource rich  
449 and/or more heterogeneous than small patches. This may make it possible for closely-related  
450 species to influence each other in small (less resource rich and more homogeneous) patches  
451 but not in larger, more heterogeneous and more resource-rich patches. This postulate would

452 be consistent with work by authors such as Jeltsch et al. (1998) who have found that fine-  
453 scale habitat heterogeneity can facilitate co-existence. Similarly, competition between species  
454 might be particularly pronounced when resources are limited (Ford 1979, Newton 1998). On  
455 this basis, a key future research task will be to carefully measure the structural and floristic  
456 variability of the woodland patches in our study area and determine if larger patches are in  
457 fact more heterogeneous than small patches. Such work, together with additional studies of  
458 the habitat requirements of bird species, would aim to determine if measures of micro-  
459 heterogeneity can help explain patterns of occurrence and/or recording frequency.

460         Some of the temperate woodland patches in our investigation were smaller (< 1 ha)  
461 than the territory size of individuals of the species we examined. Given this, it is possible  
462 these birds were using resources from multiple woodland patches or used such patches as  
463 stepping stones (see Fischer and Lindenmayer 2002). Such spatially dispersed multiple patch  
464 use may have provided opportunities for species within the same guild to co-occur.  
465 Conversely, patterns of negative co-occurrence observed for some members of particular  
466 guilds still imply that apparently competitively inferior species are using small patches less  
467 often, including as stepping stones to other patches.

#### 468 **Conservation implications**

469         Plantation forestry is increasingly recognised as a major driver of landscape change  
470 around the worldwide (Felton et al. 2010, Hulvey et al. 2013). Many plantations are being  
471 established in areas where patches of original native vegetation cover are retained (Bauhus et  
472 al. 2010). This study has revealed that these retained patches support some key elements of  
473 biota, but changes in the surrounding matrix can lead to significant changes in biodiversity,  
474 including the development of novel assemblages (Lindenmayer et al. 2008). After controlling  
475 for factors like time since plantation establishment and vegetation type, our results indicated  
476 that the occurrence and abundance of a particular species in rapid transforming plantation

477 landscapes can be influenced by the occurrence and abundance of other closely-related and  
478 functionally-similar species. Surprisingly, the majority of co-occurrence patterns were  
479 positive, suggesting that novel assemblages of typically forest and woodland-associated birds  
480 may promote the development of behaviours like mixed feeding flocks that might otherwise  
481 not have occurred. However, because some of the observed co-occurrence patterns were  
482 contingent upon patch size, we have identified potential “winners and losers” resulting from  
483 plantation-generated landscape transformation. For example, the competitively inferior  
484 Eastern Rosella (relative to the Crimson Rosella) and Buff-rumped Thornbill (relative to the  
485 Striated and Yellow-rumped Thornbills) are likely losers in small patches of woodland  
486 embedded within the maturing plantation-dominated landscape. From a conservation  
487 perspective, this may, in turn, lead to elevated local and region extinction risks that are not  
488 due to localised changes in habitat, but rather localised changes in competition among similar  
489 species (within the same guild) as reflected by patch size mediated effects on co-occurrence  
490 patterns. Such patch size mediated effects have significant implications for plantation design.  
491 This is because they highlight the need to ensure the retention and subsequent maintenance of  
492 large patches of remnant vegetation during plantation establishment (reviewed by  
493 Lindenmayer and Hobbs 2004) and thereby avoid elevated levels of competition between  
494 closely related species within small patches.

495         We argue that a better understanding of associations between species is important to  
496 prevent problems like inter-linked extinctions (Saterberg et al. 2013) such as those of  
497 particular species which result from losses of co-dependent or closely associated species  
498 (sometimes termed co-extinction cascades; Koh et al. 2004, Bascompte 2009). This is also  
499 useful to improve the ability to quantify the effects of landscape transformation and habitat  
500 fragmentation on biota (Meyer and Kalko 2008) as well as better predict the potential  
501 distributions of species (Elmendorf and Moore 2008), including species distribution patterns

502 in human-modified environments (Lindenmayer and Fischer 2006). Finally, we suggest that  
503 the kind of work reported here is important as many studies recognize the need to conserve  
504 not only species themselves, but also the associations among species as these are also critical  
505 in the functioning and assembly of ecological communities (Bascompte and Jordano 2007,  
506 Tylianakis et al. 2010).

### 507 **Concluding remarks**

508 We found that after controlling for factors like time since landscape change, vegetation  
509 type and other factors, the occurrence and abundance of a particular species in landscapes  
510 undergoing rapid transformation can be influenced by the occurrence and recording  
511 frequency of other closely-related and functionally-similar species. Co-occurrence can be  
512 either positive or negative and may also vary between different groups of species, even  
513 within a given guild. Co-occurrence also can be influenced by factors like patch size,  
514 sometimes in quite unanticipated ways. Together, our findings underscore the complexity of  
515 biotic responses to rapid large-scale landscape transformations, like plantation expansion, and  
516 indicate that some responses can be unexpected.

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643

644

645

646 **Table 1.** Number of detections of each bird species in each year. The values in a given cell  
647 correspond to the number of eucalypt woodland remnants (out of 55) in which a particular  
648 species was recorded in a given year. Note that including a lagged value of the response  
649 resulted in data from 1998 and 2000 not being incorporated in the statistical modelling.  
650

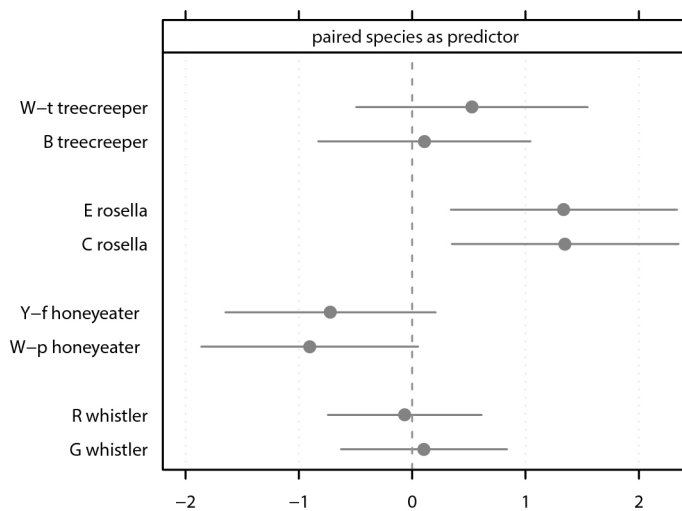
<b>Species/ Year</b>	<b>1999</b>	<b>2001</b>	<b>2003</b>	<b>2005</b>	<b>2007</b>	<b>2009</b>	<b>2011</b>
Brown Thornbill	19	16	20	31	45	49	51
Yellow-rumped Thornbill	13	11	7	6	6	10	7
Buff-rumped Thornbill	7	6	11	9	10	14	11
Striated Thornbill	12	13	11	25	24	26	36
Brown Treecreeper	8	5	4	3	5	3	3
White-throated Treecreeper	22	19	28	28	23	26	28
Crimson Rosella	27	32	35	33	39	42	41
Eastern Rosella	8	8	6	9	4	3	8
Golden Whistler	0	4	7	11	11	16	16
Rufous Whistler	29	29	40	41	39	43	42
White-plumed Honeyeater	18	28	15	15	15	9	3
Yellow-faced Honeyeater	30	36	43	54	53	51	55
Grey Fantail	32	41	40	48	49	54	55
Willie Wagtail	15	17	16	13	6	12	4

Restless Flycatcher	4	8	3	1	4	2	3
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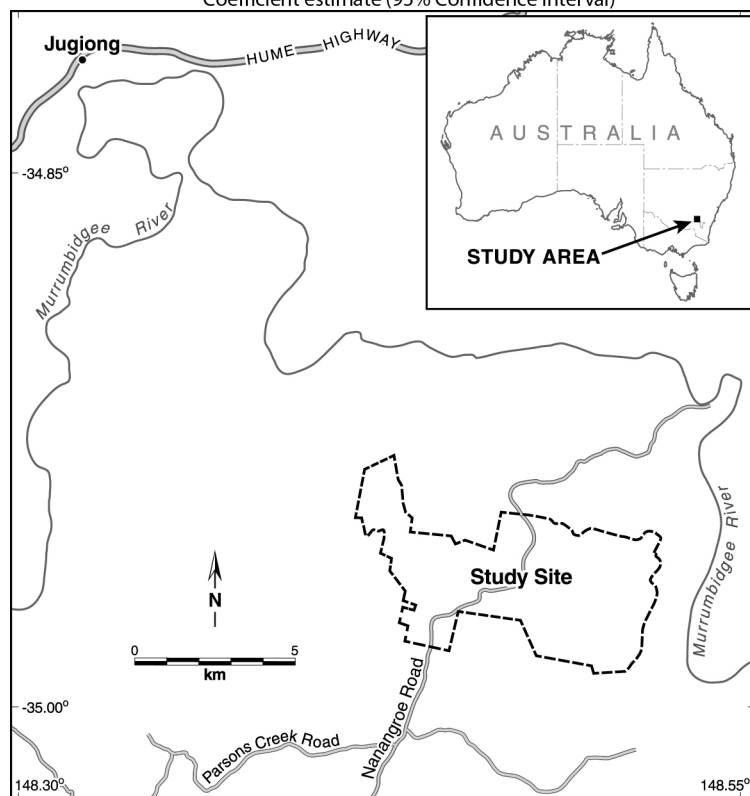
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653 **Figure 1.** Location map of Nanangroe study site

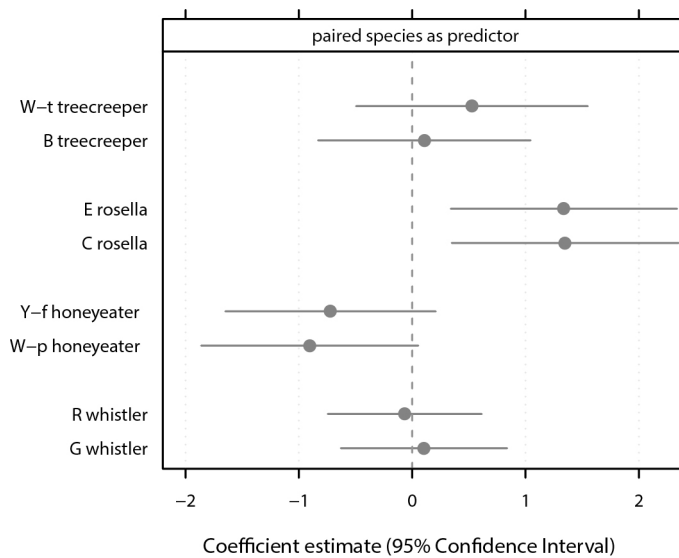


654 Coefficient estimate (95% Confidence Interval)



655  
 656 **Figure 2.** Regression co-efficient estimates for guild member species as predictor from the  
 657 occurrence part of the two-part model. Estimates are presented in pairs where the response  
 658 species is listed in the left-hand margin with the ‘paired species’ included as an indicator  
 659 variable. Co-occurrence is inferred from the model where there are significant ‘matching’  
 660 terms for the predictor species variable within a guild member pair (e.g. rosellas).

661



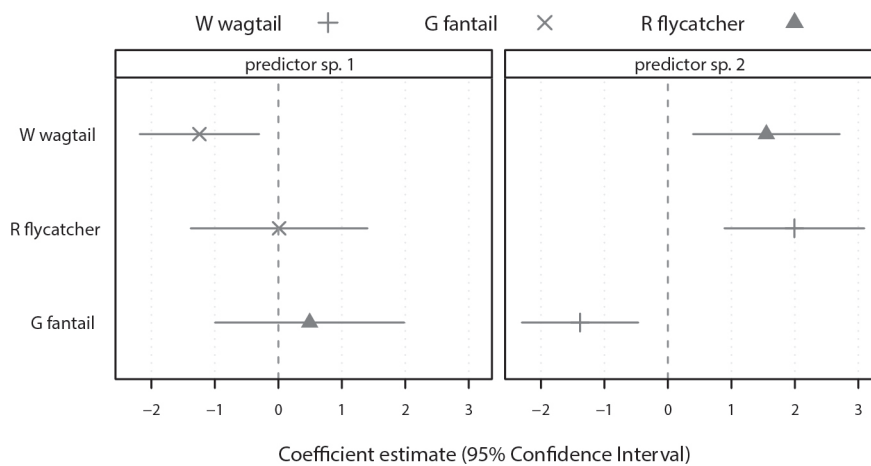
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665 **Figure 3.** Flycatcher regression coefficient estimates for guild member species as predictors  
 666 for the occurrence part of the two-part model. With three guild member species there are two  
 667 predictor species in each model. For example, negative co-occurrence or avoidance is  
 668 inferred between the wagtail and the restless flycatcher due to reciprocal, significant negative  
 669 terms for the predictor species variable for these two species.

670



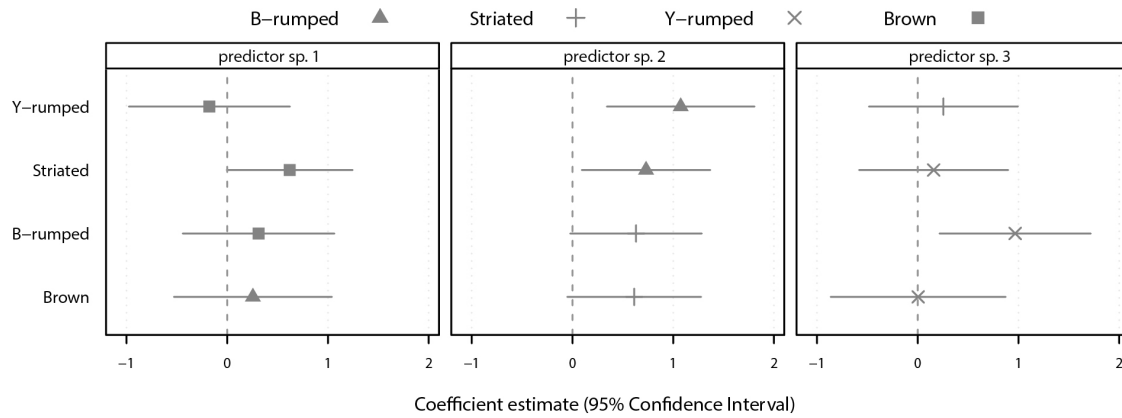
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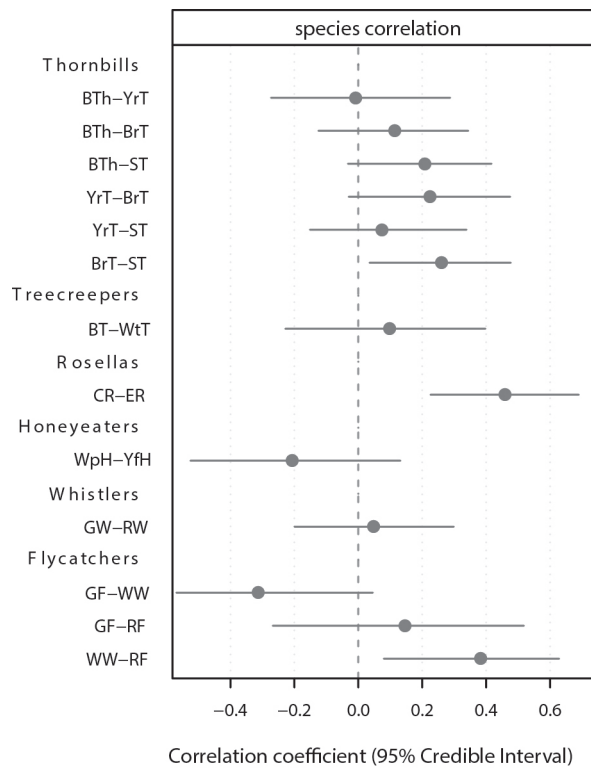
674 **Figure 4.** Thornbill regression coefficient estimates for guild member species as predictors  
 675 for the occurrence part of the two-part model. Each model has the occurrence of three guild  
 676 member species as predictors. For example, positive co-occurrence is inferred between the  
 677 Brown and Striated thornbills.



678

679

680 **Figure 5.** Correlation between paired species for the Bayesian multivariate logistic regression  
 681 models. Six models were fitted – one for each guild. Comparing the outcome of Bayesian  
 682 multivariate logistic regression modelling with the univariate two-part models showed that  
 683 the results were largely consistent between the two kinds of data analysis.



684