

Landscape scale drivers of pollinator communities may depend on land use configuration

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35 Abstract

36 Research into pollinators in managed landscapes has recently combined approaches of pollination- and landscape ecology, because key stressors are likely to interact across wide 37 38 areas. While laboratory and field experiments are valuable for furthering understanding, studies are required to investigate the interacting drivers of pollinator health and diversity 39 across a broader range of landscapes and a wider array of taxa. Here we use a network of 96 40 41 study landscapes in six topographically diverse regions of Britain, to test the combined importance of honey bee density, insecticide loadings, floral resource availability, and habitat 42 diversity to pollinator communities. We also explore the interactions between these drivers 43 and the cover and proximity of semi-natural habitat. We found that among our four drivers, 44 only honey bee density was positively related to wild pollinator abundance and diversity, and 45 the positive association between abundance and floral resources depended on insecticide 46 loadings and habitat diversity. By contrast, our exploratory models including habitat 47 composition metrics revealed a complex suite of interactive effects. These results demonstrate 48 49 that improving pollinator community composition and health is unlikely to be achieved with general resource enhancements only. Rather, local land-use context should be considered in 50 fine-tuning pollinator management and conservation. 51

- 52
- 53
- 54 Keywords
- agriculture, bumblebees, competition, hoverflies, land-use, solitary bees

57 Introduction

58 The health of insect pollinator populations and communities has become a topic of global importance in recent decades, not least because of widely reported declines [1, 2] and the 59 60 reliance of ecosystems on pollination services [3]. Pollinators are under pressure from multiple interacting stressors [4], with clear physiological and behavioural implications of 61 management practices such as insecticide application [5, 6], honey bee hive placement [7, 8] 62 and floral resource enhancement [9]. Laboratory and semi-field studies of these impacts often 63 focus on individual species such as the managed honey bee or key bumblebee species. Yet 64 attention is beginning to turn towards the importance of combining the approaches of 65 pollination ecology and landscape ecology [3, 10], particularly as the key stressors of 66 pollinator decline are likely to interact across wide areas [11]. In this study, we examine the 67 combination and interaction of landscape factors that are likely to affect pollinator 68 populations and communities across the widely varying UK countryside. 69

70

Our understanding of pollinator health has advanced significantly over the past decades by 71 studies conducted on small groups of species or at small scales. For example, exposure to 72 73 toxic pesticides can directly affect health, indirectly impact performance via foraging and reproduction [12, 13], and also impair immune responses to pathogens [4]. High densities of 74 managed honey bees may be a stressor for wild bee populations under some conditions [7], 75 due to competition for floral resources [14], or due to increased risk of pathogen spill over 76 [15]. Similarly, poor nutrition due to low quality floral resource provision can increase the 77 78 incidence of disease in honey bees [16], and pathogens are more likely to be spread in landscapes with low flower diversity [15, 17]. However, we are also discovering some 79 interactions between these and other important drivers, such as the moderating influence of 80 diverse floral resources and semi-natural habitat on the effects of agricultural chemicals on 81

insect development [18, 19]. The stressors to this diverse guild of insects are manifold and
interactive [4], yet we lack comprehensive, standardised field studies to fully demonstrate
how these factors influence the wider pollinator community [3, 20].

85

Research into landscape scale drivers of pollination populations and communities has grown 86 significantly in the last two decades [3, 10, 11, 20-22], to the extent that landscape scale 87 pollinator conservation is strongly encouraged by governments [3]. However, further research 88 is needed to synthesise the impacts of widely-diverging land management practices across 89 90 topographically diverse countries and to determine context-specific recommendations. Recent research has made great strides in identifying the important interactive effects of habitat type, 91 landscape configuration and other drivers, but studies are often only focussed on particular 92 crop types [23, 24], certain habitat types [25, 26], or limited species groups [27, 28]. 93 However, we require further studies to examine the scenarios in which landscape composition 94 95 and configuration are important in mitigating the impacts of drivers such as habitat loss and fragmentation [29-33]. This information would be of considerable use when identifying 96 landscape features to be prioritised for safeguarding pollinator communities (e.g., [34]). 97

98

In this study, we use a network of landscape study sites representing the full land use gradients in six regions of Great Britain. Our site selection protocol was designed to test the combined importance of four well-documented landscape drivers of pollinator community health (honey bee density, insecticide loadings, floral resource availability, and habitat diversity [35]). We surveyed the study sites for a wide range of pollinating insects for two years and aimed to understand how these land use factors are linked to pollinator density and diversity, which we use as proxies for community health. We predicted that, in line with

previous work, managed honey bee densities and insecticide application would have negative 106 107 impacts on wild pollinator community composition across the country [7, 36]. Conversely, we predict floral resources and habitat diversity to have positive impacts due to their importance 108 in enhancing pollinator health at individual, population and community levels [10, 26, 37]. In 109 addition, as resource provision has the potential to offset the negative effects of intensive 110 agriculture [10, 15, 18], we expected to find similar interactions between our drivers. 111 112 Furthermore, during field surveys we observed that the configuration of wide ranging habitat types are likely to play important roles in pollinator community composition in British 113 landscapes, as also shown elsewhere [38, 39]. Therefore, we also explore the potential for 114 115 these land use factors to enhance our positive drivers and mitigate negative ones [3].

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117

118 Methods

119 *Pollinator health*

In this study we use measures of pollinator community composition, including abundance and 120 diversity, as proxies for community health. There are limitations to this approach because 121 community health is typically measured across several years, requires historical baseline data 122 123 and/or involves direct measurements of fitness (e.g. longevity, reproductive success) [40]. 124 However, such data are difficult to collect over multiple landscapes and for entire 125 communities. We therefore use more convenient diversity and abundance measures as indicators of community health, as it is reasonable to expect that landscapes with many 126 127 populations able to optimally utilise resources to improve fitness, are likely to result in diverse and abundant communities. This is not always true (e.g., [41]), but wild bee abundance and 128 diversity have been correlated with pollinator success in some systems (e.g., [42]). 129

130

131 *Study site selection*

132 A detailed account of the selection of our study regions and sites is published elsewhere [35],

- but we will provide a brief overview here. We first selected six 100 x 100 km "focal regions"
- to represent the vegetation and bioclimatic gradients of Great Britain. All possible
- 135 combinations of six 100 km grid squares covering the country were measured in terms of the
- proportional area of all broad habitat types (using the 2007 Land Cover Map; [43]). The
- 137 process was repeated for the Institute of Terrestrial Ecology (ITE) land classes (a stratification
- 138 of all British 1 km squares allowing for representative, unbiased sampling given topography,
- 139 climate and human infrastructure; [44]), and the six-region combination that provided the
- 140 closest representation of Britain in both respects was selected (Fig 1).





Figure 1: Map of the six 100 x 100 km study regions (black squares) selected to represent
Britain in terms of broad habitats, topography and climate. The black dots depict the 16 study
sites chosen within each region.

Within each of the six regions, 16 study sites measuring 2 x 2 km were selected along four 146 147 gradients: 1) honey bee densities, estimated from Beebase (www.nationalbeeunit.com) database information on local colony densities, and weighted by distance using data on honey 148 bee foraging distances; 2) Insecticide loadings (including the loadings derived from 149 150 insecticidal properties of fungicides and herbicides), measured as a summed honey bee hazard score, estimated from areas of 36 crop groups and insecticides usage data from the Pesticide 151 152 Usage Survey; 3) Floral resource availability (kilograms of sugar from nectar per hectare per year), estimated by combining flowering species cover (insect-pollinated species including 153 trees and bushes) per unit cover of each habitat from the Countryside Survey 2007 [45] and 154 155 models of per-flower nectar quantity parameterised using field measurements of nectar 156 production [46]; and 4) Habitat diversity values, calculated as Shannon diversity indices using habitat cover data from the 2007 Land Cover Map [43]. Full details of these estimates can be 157 found in the Supplementary material. 158

159

160 To select the 16 sites in each region, we first scored all possible 2,500 grid squares of each region along the four gradients, standardised the values and applied a selection algorithm to 161 find the sixteen sites that maximised the difference between high and low values of the four 162 drivers and the orthogonality between them. For full details on the field site choice, see [35]. 163 The final 16 sites chosen for each region were thus considered to represent every combination 164 of relatively high and low values for each of the four gradients. The values of the gradients of 165 the final sites were subject to validation over the two-year survey period [35] and validated 166 scores are used as predictor variables in this study. Insecticide loadings were adjusted first 167 with ground referencing habitat and crop types. This resulted in a large number of our sites, 168 particularly in Scotland and northern England, having insecticide loadings corrected to zero, 169 because arable fields detected by the LCM 2007 were often reseeded grassland. Loadings for 170

sites with confirmed conventional crops were further validated via questionnaires provided to 171 172 some landowners (where land ownership could be identified). For sites with confirmed chemical applications, the correlation between estimated and validated loadings was $r_s = 0.67$ 173 [35]. Floral resource availability was validated through flower species surveys collected 174 during the studying period and statistically modelled nectar availability (see Supplementary 175 Materials for details; correlation with estimated values: $r_s = 0.28$); and habitat diversity was 176 177 validated by field surveyors confirming or correcting maps of broad habitat classes (correlation with estimated values: $r_s = 0.77$) [35]. However, we were unable to improve on 178 our original estimates of the honey bee density variable, as honey bees are poorly represented 179 180 in pan trap samples ([47]; see below), and so the original modelled estimates of this gradient 181 were retained.

182

We selected sites based on their values at the "tetrad" 2 x 2 km scale because this is the finest scale at which most datasets are available, and due to the relatively high mobility of many European pollinators [48]. However, as many solitary bees tend to forage across much smaller scales, we also calculated floral resource availability and habitat diversity for a central "inner" square (667 x 667 m) at each site (where pollinator collection was conducted; see below). We then tested whether these inner square variables were preferable predictors of pollinator responses to tetrad level variables (see Data analysis below).

190

191 *Pollinator collection*

In each region, a team of two surveyors was employed each year to collect pollinator
specimens following a standardised protocol. Pollinators were trapped using pan traps
consisting of three bowls painted yellow, white and blue with UV-reflecting paint [47], and

attached to a wooden stake at the height of the vegetation. These traps are "activity-based" 195 196 with the colours acting as an attractant to foraging insects. It is possible that local flowering plant context affects the effectiveness of such a method. We attempted to control for this 197 possibility by measuring floral resources in the area surrounding each trap (1m radius), but the 198 variable was not significant in statistical modelling (not shown) and was omitted from final 199 200 analysis. Furthermore, previous testing has found pan traps to sample pollinator communities 201 more efficiently than observational methods [47]. Five pan traps were assigned to each site, and they were placed within a central square (667 x 667 m) of each 2 x 2 km site, and using 202 the following criteria: away from potential disturbance by livestock and humans, in unshaded, 203 204 open habitats, and approximately equidistant and at least 100 m from each other.

205

Each time a trap was set up, the bowls were half filled with water and a drop of unscented 206 207 detergent was added to break the surface tension. Whenever possible, the traps were placed 208 out when forecasts predicted clear, dry conditions and left in place for 24 hours before the 209 bowls were removed and the insect material transferred to plastic bags for later mounting. The traps were sampled three times (Round 1: May, Round 2: June-July, Round 3: August-210 September; see Table S1 for precise dates), randomising the order of survey sites each time. 211 212 Due to the geographic spread of the field sites across the region, it was usually only possible to set up pan traps in four sites at a time, although in some regions with limited access to sites 213 (e.g. Inverness-shire), fewer sites were sampled in a day. Collected insect material was pinned 214 215 and mounted during the summer of collection, and specimens were identified to species by Hymettus Ltd. Taxonomic resources included [49] for hoverflies, and test keys that formed 216 217 the basis of [50] for bees and wasps.

219 Floral resource diversity, habitat composition and habitat configuration

220 In order to explore the mitigating effect of landscape context on responses to the four key drivers, we derived three further variables. First, floral resource diversity was calculated using 221 222 the flower species and nectar data used to validate the floral resource availability driver. We first estimated each flowering species' nectar provision in µg per m² for each of 28 broad 223 habitat types (the Broad habitat sub-classes listed in Table S6, Supplementary material, plus 224 225 the linear features: hedgerows, water edges, stone walls and fencelines). We then scaled this up to the landscape scale, by multiplying the values by the area of each habitat type for each 226 site. These values were summed for each species to derive their contribution to the site level 227 228 floral resource availability. We used these contributions to calculate the Shannon diversity index of floral resources for site (see Supplementary materials for full details). We preferred 229 this measure to a flower species diversity index, because it emphasises the richness and 230 evenness of nectar sources [46]. 231

232

Second, habitat composition was defined as the percentage cover of semi-natural habitat 233 (SNH) in each site, and was derived from validated land cover data described above. We 234 235 included all habitat types not classed as arable, improved grassland, urban and open water in this calculation. Therefore, our measure of SNH comprises all aspects of forest (including 236 conifer plantations), and all types of rough, low productivity grassland. Across Britain, these 237 types of habitat may be subject to varying levels of management, but in comparison to arable 238 and improved grassland, this can be considered low intensity. Further, while conifer 239 240 plantations are not typically useful foraging habitats for wild pollinators, in many of our sites, particularly in Scotland and Eastern England, the large areas of conifer are managed as nature 241 reserves and recreational areas and may represent useful nesting habitat, structural diversity 242 and corridors for movement. We also selected this measure because it provided a broad 243

gradient of data across all six regions, and because use of separate habitat class percentagecovers as individual variables led to problems with collinearity and residual heterogeneity.

246

Thirdly, habitat configuration was estimated as an index of habitat proximity following a 247 method described by [39]. On each site, 100 m buffers contoured around patch boundaries 248 were created for each SNH patch, including linear features (features found at field margins 249 250 such as hedgerows, water features and fence lines). Subsequently, the area of overlapping buffer zones was calculated and divided by the total buffer area to represent habitat proximity. 251 252 Therefore, high values of this index are likely to represent landscapes with many closely located patches of semi-natural habitat, and low values may represent sites dominated by 253 intensively managed land types or by large patches of a single SNH type. Spatial calculations 254 were conducted using QGIS (v3.10.3 [51]). 255

256

257 Data analysis

We pooled the insect pollinator data from the five pan traps at each site and across the three 258 rounds and analysed data from both years in the same models (i.e., each site was represented 259 by two years of sampling data). We used the sampled pollinators to estimate pollinator 260 261 abundance, species richness, and the inverse Simpson diversity index (1/D). We derived these 262 measures for the 'full' wild pollinator community (all hoverflies, wasps and bees except 263 honey bees), as well as separately for bumblebees (*Bombus* spp.), solitary bees (including cleptoparasitic species) and hoverflies (Diptera: Syrphidae). These community response 264 265 measures were then analysed in two ways, using a confirmatory approach to test our original hypotheses, and an exploratory approach to assess possible mediating roles of habitat cover 266 and proximity. For the confirmatory models, we fitted generalised linear mixed models 267

(GLMM) to each response with the four drivers (honey bees, insecticides, habitat diversity, 268 269 and floral resource availability) and all two-way interactions as fixed explanatory variables. 270 Higher-order interactions were excluded for clarity and to avoid complex interpretation. To improve model convergence, insecticides and floral resource availability were log-271 272 transformed, and all drivers were scaled and centred. We tested whether the "inner" scale variables for floral resource availability and habitat diversity were better predictors than those 273 274 at the tetrad scale using AIC, and by comparing residual diagnostic plots. For all response variables, the choice of variables made negligible difference to model fit ($\Delta AIC < 2$), and we 275 proceeded with the tetrad scale variables for consistency. The exploratory approach followed 276 277 the same procedure, but included the scaled and centred variables SNH, habitat proximity and 278 floral resource diversity, and all two-way interactions between them and the four main drivers. 279

280

All data analyses were performed in the R programming environment (v4.1.0 [52]). Mixed 281 282 models were fit using the *glmmTMB* package [53], and in all cases, a fixed factor for sampling year (2012/13) and a fixed integer variable for the number of pan trap bowls successfully 283 collected (out of a total of 45) were included as covariates to account for differences between 284 years and for the effect of trap bowls being disturbed by animals or passers-by, respectively. 285 Random intercepts for 'site' (n = 96) nested within 'region' (n = 6) were also specified. There 286 was no collinearity between the explanatory variables, which was checked using variance 287 inflation factors (VIF) with the *performance* package [54]. The error distribution for each 288 response variable was determined using residual diagnostic plots and tests applied using the 289 290 residual simulation methods of the DHARMa package [55]. In most cases for count data (abundance and species richness), the negative binomial distribution with quadratic 291 parameterisation ('nbinom2' family) provided the best fit to the data, although in some cases 292

(bumblebee and hoverfly species richness) the Poisson distribution provided a better fit. The 293 294 Gamma distribution with a log link was used to model the Inverse Simpson diversity index. All model (simulated) residuals were inspected visually for assumptions of linear modelling 295 (normality and homoscedasticity). Model residuals were also tested for spatial and temporal 296 autocorrelation, and the random structure adequately accounted for the clustering and 297 repeated nature of the sampling. Following model validation, 95% confidence intervals were 298 299 calculated for all model estimates using the *confint* function of the *glmmTMB* package, which computes Wald intervals by default. Significant interactions (those with confidence intervals 300 not including zero) were plotted with simple slopes, where the predicted effect of one 301 302 interacting variable is plotted for several fixed values of the second interacting variable. In most cases, we chose to keep the second interacting variable constant at the 1st, 2nd and 3rd 303 quantile values. The exception was for insecticide loadings. Our sites were relatively evenly 304 305 distributed between those with and without insecticide loadings. We therefore kept this variable constant at zero and at the median of those sites with insecticide loadings. 306

307

308 **Results**

In total, we collected 20,236 insect pollinators representing 294 species, with a greater
number of individuals and species collected in 2012 (Table 1). Most bee individuals and
species were captured in the two southernmost regions (Cambridgeshire and Wiltshire), and a
high number of hoverflies were caught in the "middle" regions of Ayrshire, Yorkshire, and
Staffordshire. The northernmost region, Inverness-shire, had the lowest numbers of
individuals and species across all groups.

- **Table 1**: Captures of pollinator individuals (and species numbers) across the six focal regions
- and for the three pollinator groups. Individuals identified only to genus were removed from

	All pollir	All pollinators		Bumblebees		Solitary bees		Hoverflies	
	2012	2013	2012	2013	2012	2013	2012	2013	
Suffolk/Cambridgeshire	1,830	1,224	474	420	884	331	342	424	
	(142)	(116)	(10)	(9)	(66)	(45)	(35)	(43)	
Gloucestershire/Wiltshire	1,568	967	154	195	515	240	845	516	
	(126)	(94)	(9)	(11)	(59)	(34)	(39)	(41)	
Staffordshire	2,402	1,636	147	427	87	133	2,159	1,048	
	(85)	(89)	(10)	(11)	(23)	(26)	(46)	(41)	
Yorkshire	1,144	658	65 (7)	78	21 (8)	69	1,055	505	
	(50)	(57)		(9)		(13)	(32)	(32)	
Ayrshire/Renfrewshire	4,961	2,885	198	308	24	31	4,731	2,496	
	(78)	(71)	(12)	(10)	(10)	(6)	(52)	(49)	
Inverness-shire	664	297	172	114	37 (9)	18	440	149	
	(60)	(45)	(9)	(9)		(6)	(37)	(23)	
Total	12,569	7,667	1,210	1,542	1,568	822	9,572	5,138	
	(240)	(205)	(17)	(16)	(86)	(67)	(89)	(84)	

the dataset when calculating species numbers.

319

320

321 *Confirmatory analysis*

322 The four target drivers as main effects in our GLMMs did not significantly affect abundance

323 of total pollinators, or of bumblebees or hoverflies when considered separately (Table S8), but

there was a positive association between managed honey bee density and solitary bee

abundance. This relationship was also present for both total pollinator and solitary bee

richness and diversity (Table S9 & S10).

327

328 We only found two significant interactions between the focal drivers. First, the effect of floral

329 resource availability on total pollinator abundance depended on insecticide loadings in the

surrounding landscape, with the positive influence of floral resource availability most

331 pronounced when loadings were absent, and the association apparently reversed at high

- loadings, although with high uncertainty (Table S8, Fig. 2a). Second, the association between
 floral resource availability and bumblebee abundance depended on habitat diversity,
 suggesting that floral resources were more beneficial to bumblebees in landscapes with
 diverse habitats (Table S8, Fig. 2b).
- 336



Figure 2: Interaction graphs for a) the abundance of total insect pollinators plotted against floral resource availability when insecticides are absent, and when insecticides are "high" (median insecticides for sites with non-zero values), and b) bumblebee abundance for three levels of habitat diversity at the 1st, 2nd and 3rd quartile. Regression lines show the predicted abundance from the GLMM (in counts) when all other predictors are held constant at mean values. Shaded areas are \pm 1 SE. See Table S8 for full model results.

- 345
- 346 *Exploratory analysis*

The exploratory models revealed several consistent interactions between focal drivers and additional variables. Honey bee density was found to interact with habitat diversity for the abundance of all pollinators (Fig. 3a) and solitary bees separately (Fig. S2, Table S11), as well as the richness of all pollinators, solitary bees and hoverflies (Fig. S2, Table S12). The

positive association between these responses and honey bee density occurred at low to 351 352 medium habitat diversity, and the opposite occurred in more diverse landscapes (as illustrated by Fig. 3a). A similar interaction was found between honey bee density and semi-natural 353 habitat proximity for some of the same responses (total pollinator abundance: Fig. 3b, Table 354 S11; Solitary bee abundance: Fig. S2, Table S11, Total richness & hoverfly richness: Fig. S3, 355 Table S12, hoverfly diversity: Fig. S4, Table S13). For example, the simple slopes of this 356 357 model term suggests that more abundant and diverse pollinator communities occur at high honey bee densities and when SNH patches are close together, but there may be a negative 358 relationship with honey bee densities in landscapes with low habitat proximity (Fig. 3b). 359

360

361 Honey bee density was also found to interact with insecticide loadings for total and hoverfly species richness (Fig 3c & d, Table S12), and with floral resource availability for solitary bee 362 richness (Fig. 3e, Table S12) and solitary bee diversity models (Fig. 3f, Table S13). At high 363 364 insecticide loadings, there was a positive association between honey bee density and both 365 total and hoverfly richness, but the opposite pattern for hoverfly richness in the absence of insecticides. In addition, honey bee densities were more strongly positively associated with 366 solitary bee richness at lower levels of floral resource availability, and with solitary bee 367 diversity at higher levels of floral resource diversity. 368

369

370





374 **Figure 3:** Interaction graphs of the significant interactive effects of landscape drivers on a) – b) the abundance of total insect pollinators, c) total species richness, d) hoverfly richness, e) 375 solitary bee richness, and f) solitary bee diversity. In graphs c) and d), insecticide loadings are 376 either absent ("none") or "high" (median insecticide loadings for sites with non-zero values). 377 In all other graphs, the 2nd predictor level is held constant at the 1st, 2nd and 3rd quartiles. 378 Regression lines show the predicted abundance, richness or diversity from the GLMM when 379 all other predictors are held constant at mean values. Shaded areas are ± 1 SE. See Tables 380 S11-S13 for interaction confidence intervals. 381

382

383 Insecticides also interacted with floral resources, habitat diversity and the amount of SNH in

the landscape, and these were mainly found for total pollinator and bumblebee abundance

(Fig. S2, Fig. 4a, Table S11) and bumblebee diversity (Fig. S4, Table S13). Firstly, total

pollinator and bumblebee abundance were positively associated with floral resource

387 availability when insecticides were absent, and show weak negative relationship when

insecticides were present (Fig. 4a). A similar pattern is also shown for the interaction between
insecticide loadings and floral resource diversity (Fig. 4b).

390

There was a contrasting interaction between insecticides and habitat diversity for total
abundance and bumblebee diversity. Habitat diversity appears to be negatively associated
with total pollinator abundance in the presence of insecticides, but positive when they were
absent (Fig. 3c, Table S11). Conversely, habitat diversity was positively related to bumblebee
diversity in landscapes where insecticides were applied, and negatively related in the absence
of insecticides (Fig. 3d, Table S13).

397

398 A similar contrasting pattern was found in relation to SNH variables. The abundance of all pollinators and hoverflies were positively related to SNH cover when insecticides were 399 present in the landscape, but negatively related in untreated landscapes (Fig. 3e & S2, Table 400 401 S11). The interaction between insecticides and habitat proximity showed the opposite pattern 402 for total pollinator diversity (Fig. 3f, Table S13). Landscapes with no insecticide applications and SNH patches in close proximity were associated with high species diversity. However, 403 404 relatively high diversity was also related to high insecticides and low connection between 405 habitat patches.



Figure 4: Interaction plots of the significant interactive effects of landscape drivers on a) – b) bumblebee abundance of total insect pollinators, c) & e) abundance of total insect pollinators, d) bumblebee diversity and f) total species diversity. In all graphs, insecticide loadings are held constant at either absent ("none") or "high" (median insecticide loadings for sites with non-zero values). Regression lines show the predicted abundance or diversity from the GLMM when all other predictors are held constant at mean values. Shaded areas are ± 1 SE. See Tables S11-13 for interaction confidence intervals.

In addition to the interactions detailed above, floral resource availability interacted with SNH
cover for a number of abundance and richness responses. These patterns were all similar,
indicating that SNH availability promoted total abundance (Fig 5a, Table S11) and richness
(Fig S3, Table S12), bumblebee abundance (Fig. S2, Table S11) and the abundance and
richness of hoverflies when floral resources were scarce (Figs. S2 & S3, Tables S11 & S12).
Similarly, floral resource availability was important to these responses when SNH cover was
low. Surprisingly, a combination of both high SNH cover and high floral resource availability

- lead to some of the lowest number of predicted species. Finally, in addition to the interactions
 involving habitat diversity above, bumblebee diversity was predicted to be highest when both
 habitat diversity and proximity were high (Fig. 5b, Table S13).
- 426



428

Figure 5: Interaction plots of selected significant interactive effects of landscape drivers on a) abundance of total insect pollinators, and b) bumble diversity. In all both graphs, the 2^{nd} predictor level is held constant at the 1^{st} , 2^{nd} and 3^{rd} quartiles. Regression lines show the predicted abundance or diversity from the GLMM when all other predictors are held constant at mean values. Shaded areas are ± 1 SE. See Tables S11 and S13 for interaction confidence intervals.

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436 Discussion
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In this study we have used the most comprehensive nationwide network of study sites toexplore the multiple, interacting drivers of insect pollinator communities in Great Britain. We

found that four landscape scale factors considered important to pollinators could not provide 440 441 simple explanations for abundance, richness or diversity patterns, except for an unexpected positive relationship between honey bee density and both total pollinators and solitary bees. 442 While some factors combined to explain total and bumblebee abundance and richness in our 443 confirmatory models, we revealed a complex set of responses when incorporating landscape 444 composition and proximity variables in our exploratory models, supporting previous work 445 446 suggesting that improving pollinator population and community health requires an understanding of local and regional land use factors [7, 10, 31]. We should note that our 447 project was not designed to test a priori hypotheses about these interactions, and we 448 449 recommend that further studies seek to confirm these findings. Caution should also be used 450 when interpreting the results involving two of our four drivers: honey bee density is the one variable that we could not validate with collected data, and the insecticide loadings variable 451 452 did not consist of as wide a range of values within regions as we would have hoped [35].

453

454 *Managed honey bee density*

We expected the estimated density of managed honey bees to have a negative relationship 455 456 with most functional groups of wild pollinators via either competition for food or the transmission of pathogens [7]. As a simple effect there were only positive associations of 457 honey bee density on total richness and diversity, and on all three solitary bee responses. 458 These may be geographical artefacts, however, because solitary bee abundance and diversity 459 generally decreased with latitude, and our knowledge of honey bee densities was likely to be 460 461 more accurate in the southern regions. It is also possible that the scale of our study sites were inappropriate for solitary bees, although finer scale measures of floral resource availability 462 and habitat diversity did not improve our models. Nevertheless, the exploratory models also 463 suggest that the positive relationship with solitary bee diversity was strongest in landscapes 464

with high floral resource availability and diversity. We expected honey bees to compete with 465 466 wild pollinators mainly for floral resources, since honey bee nesting is provided by beekeepers. However, competition is thought to be context dependent and the majority of 467 reported negative impacts of honey bees are from territories where the species is not native [7, 468 469 26]. Apis mellifera is native to the UK [56], which could contribute to the ability of honey bees and wild pollinators to co-exist in suitable locations when resources are abundant [7, 42]. 470 471 The placement of honey bee colonies in resource rich environments with coincidental healthy 472 wild pollinator communities seems an unlikely explanation, given that very few beekeepers in the UK move their honey bee colonies, instead tending to keep bees close to where they live. 473

474

We also found context dependence in the association, with only weak negative relationships 475 with abundance in landscapes with low habitat proximity (dominated by arable, grassland or 476 477 large single patches of SNH), and high habitat diversity. Sites in our study with this 478 combination of landscape properties are those with many small isolated patches of semi-479 natural habitat, which may provide nesting sites for bees, but require them to forage far into the agricultural matrix where floral resources may be scarce. Similar patterns have been 480 481 reported in Sweden, where competition between honey bees and wild bees and hoverflies increased with crop field size [57], or when the amount of semi-natural grassland in the 482 surrounding landscape was low [58]. By contrast, sites in our study with low habitat diversity 483 but high proximity to SNH, which seem to promote coexistence between managed and wild 484 pollinators are those with several large patches dominated by a single use such as moorland, 485 rough grassland or even improved grassland, and divided by linear features (e.g. hedgerows, 486 487 ditches and fence lines). These sites may be ideal situations with abundant resources for both managed honey bee hives and pollinator communities, and we suggest focussed research on 488 these large habitat types. Interestingly, there remain significant gaps in our understanding of 489

490 how honey bees can influence population-level responses of plant communities, such as plant491 abundance or distribution [7], perhaps masking wider benefits to wild pollinators.

492

493 Insecticide loadings

494 This variable was less well distributed among the regions because virtually no insecticides 495 were applied in the three northernmost landscapes, but large amounts were applied in the two 496 southernmost regions [35]. Nevertheless, in landscapes where insecticides were not applied, 497 we found positive effects of other drivers such as floral resource availability, resource 498 diversity and habitat proximity. Interestingly, floral resource availability and diversity appeared to have a negative association with pollinators in the presence of insecticides. We 499 interpret this as an increased exposure to insecticide in the presence of abundant and diverse 500 food resources, such as in chemically treated mass-flowering crops [59], or because forage 501 plants in adjacent uncultivated habitats can be sources of insecticide exposure for pollinators 502 503 via drift or soil pathways [36, 60, 61]. While field microcosm experiments suggest that diverse forage sources provided as alternatives to mass-flowering crops should offset the 504 negative impacts of insecticides [18], we did not find evidence of this at the landscape scale. 505

506

There was also a positive relationship between SNH and total pollinator abundance, and between bumblebee diversity and habitat diversity in the presence of insecticides, in line with previous findings [19, 62]. These studies suggest that higher amounts of SNH in the landscape support pollinator communities by providing a refuge from intensive agricultural practices such as chemical applications [19], although pesticide residues have been found in sites with up to 89% semi-natural grassland in the surroundings [63]. It is not clear why SNH cover had a negative impact in the absence of insecticide application, but may be due to our inclusion of

conifer plantations in SNH. Untreated sites with high SNH are likely to be those in the 514 515 northern regions with high covers of conifer and moorland, and these sites typically had low pollinator catches. Exclusion of conifer plantations from our SNH variable would have 516 resulted in these sites scoring very low on the SNH scale, perhaps nullifying the interaction 517 effects shown here. In any case, as our study design was limited in detecting within-region 518 519 relationships between insecticides and pollinators [35], future work on this scale should base 520 landscape selection on ground-truthed chemical application data. For example, while the chemical application data we used was of a high standard, our reliance on the LCM 2007 to 521 select sites with "high" estimated insecticides prevented this gradient from reflecting the full 522 523 range of loadings in Britain. Further studies could also focus on the indirect effect of herbicides via floral resources. 524

525

526 Floral resource availability

527 As well as a positive relationship with pollinator abundance and diversity in the absence of insecticides, floral resources were important to bumblebee abundance in landscapes with high 528 habitat diversity. This is unsurprising as a diversity of habitats provides a range of nesting 529 530 resources for bumblebees [26], and a correspondingly high level of continuous food supply is required to support healthy colonies [64]. As central place foragers, bees are more likely to 531 forage efficiently when flowering plants are abundant within a short distance of the nest [65]. 532 A more surprising result is that we did not find the same synergistic interaction for more 533 groups. This is perhaps because our scale of study was not appropriate for solitary bees with 534 shorter foraging ranges, for example. 535

In contrast to the above pattern, we found that floral resources were important in landscapes 537 538 with low cover of semi-natural habitat. This supports theories of floral provision in agricultural landscapes, where small patches of nesting resources, such as semi-natural 539 habitat, should be interspersed with rich floral resources to benefit pollinators [20, 31]. In our 540 541 landscapes, the combination of low SNH cover and high floral resource availability corresponds to sites with high arable cover including mass flowering crops or with a high 542 543 cover of improved grassland with flower rich field boundaries. Bumblebees may be particularly attracted to mass flowering areas over SNH [38, 65], and other bees and 544 hoverflies may benefit from the connectivity effect provided by floral resources in field 545 546 margins [20]. Conversely, the apparent negative relationship between pollinators and SNH 547 under high floral resources occurs in sites with large areas of heathland and rough grazing. In such wide, open places, pollinators may concentrate around patches of flowers rather than 548 disperse [66] and are likely only attracted to our pan trap bowls when resources are low. 549 Alternatively, floral resources may be relatively homogenous at these sites resulting in low 550 abundances of pollinating insects [30]. 551

552

553 Habitat diversity

We expected habitat diversity in general to have positive associations with the diversity of the 554 pollinator community, as a greater array of habitat cover types provide a range of alternative 555 nesting substrates and niches [26]. However, as we have shown, this can be mediated by 556 landscape context such as local honey bee densities and insecticide loadings. Furthermore, in 557 558 our landscapes low habitat diversity can correspond to large covers of intensive land uses such as arable or improved grassland, or conversely to a dominance of SNH such as heathland 559 or low intensity habitat such as coniferous forest. When other habitat variables were included 560 in models, habitat diversity showed the expected positive relationship with bumblebee 561

diversity when habitat proximity was also high. This supports findings that the provision of habitat patches *per se* is not always sufficient to promote all aspects of pollinator community abundance and diversity, but that habitat patches should be connected or at least within foraging range of a variety of functional groups [20, 31].

566

567 Conclusions

Our results are difficult to distil into simple, generalisable statements. We found rather few 568 569 simple effects of the often-cited key drivers of pollinator community composition and 570 distribution across highly variable topographic areas. This suggests that such variables do not generalise well across regions that are characterised by their land use, climate and 571 572 management. While we have not directly measured pollinator fitness, we infer from these results that improving pollinator community health at the landscape scale is also unlikely to 573 have a quick or general fix. When it comes to conservation or restoration of pollinator 574 575 communities, our study supports others studies that call for taxon- and context-specific decisions to be made [7, 32, 67]. Furthermore, unlike other studies that find no effect of SNH 576 on pollinator communities [29, 68, 69], we find support for studies that include landscape 577 578 composition and configuration variables as interactive terms in models [38, 70]. As the reality of interacting landscape drivers and their effects on pollinator community composition and 579 health is likely to be even more complex than what we have been able to test, we further 580 recommend that better policy and practice decisions are likely to be reached by taking multi-581 582 driver, multi-taxa approaches.

583

584 Despite the complexity of our results, some key messages are clear. First, pollinator
585 community health, if it is indeed correlated with abundant and diverse pollinator assemblages,

is likely to be enhanced by increasing the availability and diversity of floral resources, but the 586 587 landscape context in terms of insecticide loadings, habitat diversity and habitat proximity should be considered in their selection and placement. Second, in intensively managed 588 landscapes, floral resources can be important when SNH cover or proximity is low, and 589 habitat diversity and configuration can also play important, though complex roles. Third, 590 while other studies have found that beneficial resources can offset negative influences, we did 591 592 not find consistent evidence of this. Thus, instead of simply relying on boosting pollinator resources to rectify otherwise unhealthy management practices, we recommend that pollinator 593 conservation should be fine-tuned in relation local land use context. Finally, we re-iterate that 594 595 many of our findings were revealed from exploratory data analysis, and we did not have sufficient data for cross-validation. We therefore further recommend future landscape scale 596 research confirming the importance of habitat context to the drivers of pollinator 597 598 communities.

599

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