1	Running headline: Compositional and functional indices for biodiversity monitoring.
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3	Using compositional and functional indices for biodiversity conservation monitoring of
4	semi-natural grasslands
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6	Rob J. Lewis <sup>a,c*</sup> , Robin J. Pakeman <sup>a,</sup> Stewart Angus <sup>b</sup> and Rob H. Marrs <sup>c</sup>
7	
8	<sup>a</sup> James Hutton Institute, Craigiebuckler, Aberdeen, AB15 8QH, UK; <sup>b</sup> Scottish Natural
9	Heritage, Great Glen House, Leachkin Road, Inverness, IV3 8NW, UK; School of
10	Environmental Sciences, University of Liverpool, Liverpool L69 3GP, UK.
11	*Corresponding author: University of Tartu, Institute of Ecology and Earth Science, Lai 40,
12	Tartu, 51005, Estonia.
13	Email: robert.lewis@ut.ee
14	
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## 18 ABSTRACT

19 Compositional indicators (i.e. indices that focus on the identity of species, genes or phylogeny) have been widely used to estimate and monitor biodiversity, however, their use in 20 21 combination with species and/orcommunity functional characteristics remains limited. Using 22 large-scale, spatio-temporal data, we use both compositional and functional indices to 23 investigate land-use change impacts on the vegetation of a semi-natural grassland ecosystem 24 (Machair) for fourteen regions in Scotland, UK. Our study aimed to identify national- and 25 regional-scale temporal vegetation patterns, and through use of simple compositional and 26 functional indices (e.g. Competitor, Stress, Ruderal and Ellenberg scores) link observed 27 changes to agricultural intensification and/or land-use abandonment. Using linear-mixed 28 modelling and nonmetric multi-dimensional scaling, we showed significant national and 29 regional-scale changes in species composition over time. Increases in diversity, particularly 30 gains in Machair grassland, identified several regions that may have benefited from past 31 government incentivised schemes to protect the Machair, but which may also be suffering from 32 an extinction lag. Shifts in plant functional signatures (CSR & Ellenberg values) identified 33 varying degrees of internal (competition) and external (land-use) factors, highlighting several 34 regions where biodiversity change could be linked to reduced disturbance (i.e. lower grazing 35 intensity) or greater disturbance (i.e. land-use intensification). Our results demonstrate the 36 utility of simple compositional and functional indices for monitoring biodiversity of semi-37 natural grasslands and identifying land-use drivers of change across different spatial scales.

## 38 **1. INTRODUCTION**

39 Semi-natural grasslands (SNGs) constitute one of the most species-diverse communities 40 in the World at small spatial-scales (Wilson et al. 2012). In Europe, the high level of 41 biodiversity, characteristic of SNGs, is closely linked to century long associations with 42 agricultural land-use (Eriksson et al. 2002; Poschlod and WallisDeVries 2002). However, for 43 these tightly coupled social-ecological systems, cessation of traditional management (e.g. 44 extensive grazing, hay making and rotational arable agriculture) in favour of land-use 45 intensification or complete cessation of any form of management has become a widespread 46 occurrence (refs). Typically linked to societal changes in response to economic factors, such 47 land-use changes are considered the principal cause of habitat deterioration, fragmentation and 48 biodiversity loss among European agricultural landscapes (Benton et al. 2003; Lindborg et al. 49 2008). Therefore, given current and forecasted global biodiversity loss (Sala et al. 2000), 50 appropriate design and implementation of conservation management for attempts to maintain 51 and enhance these systems has global as well as local and regional importance. 52 However, there is a growing concern regarding the effectiveness of current monitoring 53 frameworks for attempts to conserve and enhance biological diversity (Lindenmayer and 54 Likens 2010), particularly of semi-natural landscapes (de Bello et al. 2010). For semi-natural 55 grasslands, one part of the problem is likely to be the broad spatial grain common to many 56 large-scale monitoring studies (e.g. The Countryside Survey of Great Britain; http:// www.countrysidesurvey.org.uk/). European SNGs typically exhibit high spatial heterogeneity, 57 58 usually as a result of spatially and temporally structured variation in agricultural management 59 (Plieninger et al. 2006). Moreover, variation is intensified where land is apportioned to 60 multiple landowners. Although such heterogeneity directly benefits the biodiversity of SNGs 61 (Öster et al. 2007) and surrounding agricultural landscapes (Blitzer et al 2012), patterns of 62 biodiversity change as a result of changing land-use (i.e. intensification vs abandonment) 63 become extremely variable across relatively small spatial scales. As a result, management and

conservation policies designed and applied across large spatial extents may not always be
appropriate (Fortin and Dale 2005), often failing to be successfully translated across highly
heterogeneous landscapes (Stenseke 2006).

67 Another part of the problem is the complexity of biological diversity and the suite of indicators designed to measure its various facets. This issue was recently addressed by de Bello 68 69 et al. (2010) who derived a framework to help standardise and converge the use of biotic 70 indicators (i.e. direct measures of components of biodiversity) and biotic drivers (i.e. factors 71 that govern local biodiversity) for monitoring biodiversity change, specific to semi-natural 72 habitats. Biotic indices (e.g. species richness, Shannon diversity, Simpson diversity, see 73 Magurran 2004) are commonly applied to measures of alpha diversity (sensu Whittaker 1960, 74 1972) or a subset of alpha, targeted at indicator species (i.e. suites of species that commonly 75 occur in a habitat under different environmental conditions – e.g. grazed or un-grazed, 76 favourable or non-favourable conservation condition etc; see Rosenthal 2003; Wittig et al. 77 2006). Such compositional measures are relatively easy to calculate and understand (Lamb et 78 al. 2009) but they fail to provide information of potential abiotic or biotic drivers governing 79 observed changes. On the other hand, direct measures of biotic drivers (e.g. pH, organic matter 80 and nitrogen content), for which links with biodiversity have been well researched (Hartley et 81 al. 2003; Pärtel et al 2004), are laborious and cost intensive, and therefore often unavailable. 82 This is especially true for large-scale spatial temporal monitoring studies.

To overcome these limitations, one approach is to use biotic indicators as surrogates for biotic drivers, particularly those that focus on the functional characteristics of species and ecological communities (i.e. functional indicators *sensu* de Bello et al. 2010). It is well understood that plant functional characteristics, pre-adapted to the local environment, provide distinct signatures to drivers of change that are not always easily measurable, and which often relate well to land-use processes (Pakeman 2004; Quetier et al. 2007; Lewis et al. 2013). For example: the Competitor-Stress-Ruderal (CSR) classification of plant functional types (Grime

90 1974) is frequently used to describe variation in vegetation through its functional composition 91 (Cerabolini et al. 2010; Bornhofen et al. 2011). Among SNGs, shifts in CSR functional 92 strategies can be observed. Land-use abandonment leads to size asymmetric competition for 93 light (Mason et al 2011; Laliberte et al 2013), resulting in competitive displacement of grazing 94 tolerant specialists (Tilman and Lehman 2001). Alternatively, declines in stress tolerant plant 95 species should be observed with soil nutrient enrichment brought through land-use 96 intensification (Maskell et al. 2010). Similarly, Ellenberg indicator values for plants (Ellenberg 97 1988), which define species habitat preferences and distribution, provide clear linkages 98 between plant species occurrences and the environment, proving valuable for detecting 99 changes in species compositions as a result of changes in land-use management. For example: 100 Ellenberg N scores prove to be good correlates of mineralisable nitrogen (Rowe et al. 2011) 101 and grassland productivity (Wagner et al. 2007).

Understanding the nature of vegetation change over time in this way can be paramount
to the appropriate design of nature conservation practices, policies and land management
across local, regional and national scales for all terrestrial habitats, not least SNGs. Coupled
with monitoring indicator species, functional classifications such as plant CSR strategies,
and/or Ellenberg values can prove extremely useful for highlighting the effects of land-use
impacts.

108 In this study we demonstrate the potential utility of simple functional indicators as 109 proxies of biodiversity drivers in indentifying temporal shifts in land use patterns among 110 SNGs. Using archived biological records and revisitation data, valuable for quantifying long-111 term patterns of vegetation change (e.g. McCollin et al. 2000; Bennie et al. 2006; McGovern et 112 al. 2011), we quantify national- and regional-scale temporal shifts in plant species diversity and 113 composition for an internationally important SNG (machair). Through utilising simple land-use 114 measures and functional indicators, surrogates for abiotic/biotic drivers, we link changes in the 115 vegetation to changes in land-use, and identify regions where machair SNGs are subject to

land-use intensification and those that are subject to management declines over the last three
decades. We highlight regions of potential conservation concern and discuss how simple
environmental surrogates can be a useful, cost effective measure for multi-scale biodiversity
monitoring.

120

#### 121 **2. METHODS**

122 **2.1 Study Area** 

123 We used the grassland vegetation of the Scottish Machair as a case study (Fig. 1). 124 Confined globally to the northwestern fringes of Europe, sampled were14 regions, 125 encompassing almost the entire national resource of Scottish machair, i.e. over two thirds of 126 the World's total machair extent (ca. 30,000ha; Dargie 2000). Machair SNGs make a good 127 study system as their high floristic diversity and conservation value is tightly coupled to 128 century-long associations with human settlement (Gilbertson et al. 1996) and land-use 129 practices (Supporting Information S1), and presents a clear example of how SNGs are 130 threatened through agricultural and socio-economic change (Pakeman et al. 2012).

131

#### 132 **2.2 Floristic Data**

133 Species compositional data were collated from two sample periods. Baseline data was 134 taken from the Scottish Coastal Survey (1975-1977; Shaw et al. 1983). The temporal aspect 135 was completed through a partial re-survey between 2009 and 2010 of regions known to include 136 machair communities, i.e. all transient communities from seaward embryo dunes to inland sand 137 affected peatlands, including machair grasslands (Angus 2006; Supporting Information S2). 138 Re-survey methodology followed closely that of the original survey (for details see Shaw et al. 139 1983), estimating cover of all higher plants and measuring environmental proxies for land-use 140 management within 5 m x 5 m plots.

141	Data specific to the focal community (i.e. machair grassland) was abstracted from the			
142	temporal dataset. Only those plots with a National Vegetation Classification (NVC)			
143	representative of machair grassland (Angus 2006; Table S1) were analysed. The resulting			
144	matrix (Matrix 1) comprised 321 species from 853 plots, from two sample periods. Matrix 1			
145	was used to quantify biodiversity change. To reduce noise within the dataset (see Gauch 1982)			
146	Matrix 1 was further constrained, removing species present in less than 5 plots. This species			
147	data matrix (Matrix 2) comprised 213 species from 853 plots, from two sample periods and			
148	was used to quantify compositional change. A schematic representation of how the data was			
149	handled prior to analyses is provided in Supporting Information (Fig. S1).			
150				
151	2.3 Environmental Variables			
152	Environmental proxies for land-use management included measures of past and present			
153	management, current grazing intensity, presence/absence of grazing animals and categorical			
154	measures of sward height for grasses and herbs (Table 1). Methods on how grazing pressure			
155	levels and past cultivation were determined are detailed in Supporting Information (S3).			
156				
157	2.4 Statistical Analysis			
158	All statistical analyses were carried out using the R statistical software (Version 2.15.0,			
159	R Development Core Team 2012). Where necessary, data were transformed to meet			
160	assumptions of normality and homogeneity of variance.			
161				
162	2.4.1 Quantifying temporal change in biodiversity			
163	Two biotic diversity indices: i) richness (i.e. the count of all species per 5m x 5m plot)			
164	and ii) the reciprocal of the Simpson's index $(1/D)$ were calculated from Matrix 1. Paired <i>t</i> -			
165	tests, and General Linear Mixed Models (GLMM) with a poisson link function, were used to			

166 identify statistically significant national- and regional-scale shifts respectively. The GLMM 167 (M1) took the following structure: 168 (*response* ~ *year/region*, random = land-owner/plot) (M1) 169 Where, 'year' and 'region' were nested fixed factors, and land-owner and individual plots 170 formed nested random factors accounting for the expected spatial and temporal auto-171 correlation. 172 These analyses were extended to investigate subsets of species, indicative of different 173 vegetation conditions of the machairs taken from published guidance for conservation 174 monitoring (Joint Nature Conservation Committee 2004; Table S2). Selected were four species 175 indicator groups, indicative of good and poor quality machair grassland, fallow areas 176 (indicative of continuous rotational arable agriculture) and active cultivation. Statistically 177 significant shifts in mean species richness was tested using GLMM with a poisson link 178 function. Similarly, temporal change in relative abundance was also investigated using linear 179 mixed models with a residual maximum likelihood (REML) estimating procedure. Model 180 structures equalled M1. 181 182 2.4.2 Quantifying temporal change in species composition 183 Non-metric Multi-Dimensional Scaling (NMDS; Legendre and Legendre 1998) of plots 184 in two-dimensional ordination space was applied to Matrix 2, following the procedure 185 recommended by Minchin (1987). The scores of the resulting NMDS axes were subsequently 186 used as numeric values representing community composition in consecutive analyses (Kahmen 187 et al. 2005). The Bray-Curtis (Bray and Curtis 1957) dissimilarity index was used for the 188 computation of NMDS. Species contributing most to the overall Bray-Curtis measured

temporal dissimilarity were identified through similarity percentage analyses (SIMPER; Clarke

- 190 1993). NMDS was performed using the function 'metaMDS' within the *vegan* package
- 191 (Oksanen et al. 2013).

Temporal change in species compositions was measured by calculating the paired
distance between NMDS site scores for each plot for each survey year, in two-dimensional
ordination space. The statistical significance of these shifts provides a reliable indication of
species compositional change: paired t-test (national-scale) and linear-mixed modelling with
REML (M1; regional-scale).

197

#### 198 2.4.3 Linking environmental surrogates to biodiversity and compositional change

199 Environmental surrogates, proxies for land-sue change (Table 1), and measured 200 functional indicators were projected onto the NMDS ordination, linking national and regional 201 compositional change with changing land-use parameters. Significant national and regional 202 shifts between survey years for the two functional classifications: CSR and Ellenberg scores 203 were tested using paired t-test (national-scale) and linear-mixed modelling with REML (M1; 204 regional-scale). Species abundances' (Matrix 2) were summarised as compositions of these 205 functional classifications (McGovern et al. 2011), acting as environmental surrogates for 206 various land-use changes. CSR scores of Grime (1974), derived using a spreadsheet based tool, 207 described in Hunt et al. (2004) and freely downloadable, were used to provide an indication of 208 external factors likely to alter plant growth (e.g. stress through lack of plant available resources 209 and disturbance). Similarly, Ellenberg indicator values, cover weighted and adjusted for British 210 plants, taken from Hill et al. (1999), were used to reflect plant species preferences in terms of 211 nutrient level (EbN), acidity (EbR), moisture (EbW), light (EbL) and salinity (EbS). 212 Land-use disturbance surrogates (Table 1) were examined to see how well they

- 213 explained the temporal compositional shifts. Between year differences in land-use measures
- 214 were regressed against between year differences in NMDS scores through further GLMMs:

(*response* ~ *Land-use*, random = land-owner/plot)

- 215
- 216

#### 217 **3. RESULTS**

9

(M2)

## 218 **3.1. Biodiversity change**

219 National gains in biodiversity, measured through both mean species richness (SR) and 220 Simpson's diversity (Sim) were significant (Table 2). Regional-scale analyses identified 221 significant gains in SR and Sim for two regions (Colonsay and Tiree), and significant declines 222 in SR and Sim for three regions (Benbecula, Harris and Monachs. Significant gains in SR 223 coupled with significant declines in Sim were observed for two regions (Barra and S.Uist). 224 Analyses of species subsets indicative of different vegetative states identified 225 significant national gains in richness and cover of machair grassland specialists, and a 226 significant national decline in relative abundance of species indicative of arable agriculture 227 (Fig. 2). Regional-scale analyses differentiated between regions in favourable condition, where 228 significant temporal gains in machair specialists occurred (Barra, Coll, Colonsay, Islay, S. Uist 229 and Tiree), and those of conservation concern where significant temporal declines in richness 230 and cover of species indicative of good quality machair grassland occurred (Benbecula, 231 Monachs and Shetland; Fig.2). Significant declines in the number of species indicative of 232 arable and rotational arable agriculture occurred among 5 regions (Barra, Harris, Monachs, 233 North Coast and North Usit), with significant declines in cover for 7 regions (Barra, 234 Benbecula, Islay, Monachs, North Uist, Shetland and South Uist).

235

## 236 **3.2 Compositional change**

The NMDS ordination showed clear shifts in species composition between survey years (Fig. 3). This was supported by statistically significant temporal shifts in ordination scores at both national- and regional-scales (Fig 4). Projection of environmental proxies identified a positive shift along NMDS Axis 1 to indicate plant compositional change towards shorter grassy and herbaceous swards. In addition, functional classifications Ellenberg Water and Ellenberg Reaction were positively ranked with NMDS Axis 1 indicative of communities' subject to higher saturation and more acidic substrate. Regional scale analyses identified five

244 regions to display significant positive shifts (Coll, Colonsay, Islay, S.Uist and Tiree) and one 245 region to display a significant negative shift (Benbecula) along NMDS Axis 1 (Fig. 4). 246 Proxies for land-use disturbance, e.g. vegetation sward heights, evidence of livestock 247 and cultivation, as well as stress tolerant species, were all strong correlates with NMDS Axis 2 248 (Fig. 3). Patterns indicate communities positively ranked with Axis 2 to have associations with 249 cultivation practice, be it, past or present, and or low intensity rotational arable agriculture or 250 larger-scale farming. Higher Ellenberg N values were also positively ranked with Axis 2. Only 251 one region (Shetland) was found to positively and significantly shift along Axis 2 (Fig 4). 252 An opposite trend, where communities were negatively ranked with Axis 2 displayed 253 strong associations with proxies for disturbance through grazing (i.e. shorter vegetation sward 254 heights, heavy and moderate gazing, presence of sheep and species with higher stress tolerance 255 (Figs 3). Significant negative shifts along this Axis were recorded for 2 regions (Barra and 256 Coll; Fig. 4).

257

## 258 **3.3. Linking environmental surrogates to temporal change**

259 Significant national and regional shifts were indentified among the functional 260 indicators. At the national-scale, a significant increase in functionally competitive species (p < p261 (0.001) with a concomitant significant decline in ruderal species (p = 0.005) was observed (Fig. 262 5i). Eight regions mirrored this national trend, while only two (Barra and North Uist) were 263 found to have statistically significant gains and declines in competitor and ruderal species 264 respectively (Fig. 5i). Patterns not borne at the national-scale were revealed at the regional 265 level. Three of the four regions (Barra, Monachs and North Uist) displaying significant 266 declines in ruderal species were met by significant increases in stress tolerators. The opposing 267 pattern where ruderals increased and stress tolerators declined was observed for five regions, 268 with statistically significant change identified for two (Islay and Shetland). All significant 269 changes in competitors were positive.

270	Significant national and regional shifts in Ellenberg values were also observed (Fig.
271	5ii). At the national-scale community composition shifted significantly towards species
272	favouring acidic substrate (EbR, $p < 0.001$ ) and mineralisable nitrogen (EbN, $p = 0.008$ ), while
273	species favouring moist soil conditions significantly declined (EbW, $p < 0.001$ ). At the
274	regional-scale, significant changes were observed among 11 of the 14 regions surveyed (Fig.
275	5ii). Significant regional trends for EbR, EbN and EbW were consistent with the national
276	changes observed, the only exception being a significant declines in species favouring
277	mineralisable nitrogen for one region (Lewis). A general decline was observed in light
278	competitive species (EbL), with the exception of one region (Monachs) where EbL
279	significantly increased ( $p < 0.001$ ). Significant shifts among salt tolerant species (EbS) varied
280	across regions, significantly increasing (Harris, Monachs and Sanday), and significantly
281	declining (Lewis, North Coast and Tiree).
282	Temporal change in measured land-use proxies indicative of arable agriculture
283	(RecCult & OldCult) displayed a significant positive relationship with temporal compositional
284	change measured as the change in NMDS scores for both Axis 1 and 2 (Table 3). Change in
285	vegetation height classifications for graminoids over 50cm tall significantly and negatively
286	related to compositional changes measured as the change in NMDS scores for Axis 1.
287	Temporal change in the herb layers classifications i.e. Hrb<20 & Hrb>50 were also shown to
288	have a significant positive relationship with species compositional change for Axis 1 and Axis
289	2 respectively (Table 3).
290	

## 291 4. DISCUSSION

## **4.1. Biodiversity change: linking shifts to species indicator groups**

The biodiversity value of semi-natural grasslands is well understood to be promoted by
low-intensity land-use management (Eriksson et al. 2002; Poschlod and WallisDeVries 2002).
For machair grasslands, century-long associations with human land-use (Gilbertson et al. 1996)

296 have resulted in these habitats developing in tandem, with a constant disturbance regime that 297 can now be considered a 'natural part' of the local habitat conditions. Given this tight historical 298 coupling between the biodiversity value and land-use practices, temporal changes in 299 biodiversity are likely to be linked to changes in the form, frequency and intensity of land-use 300 disturbances. In this study, we report significant national and regional temporal changes in 301 plant species diversity of Scottish machair grasslands. For many regions biodiversity significantly increased, coupled with increases in richness and relative abundance of machair 302 303 grassland specialists. These positive changes in diversity were, for many regions linked to 304 concomitant gains in plant species indicative of rotational arable agriculture. Collectively these 305 species can be viewed as favourable machair species, i.e. species that belong to the habitat 306 specific species pool. A high proportion of these species can suggest a community to be more 307 complete, relative to the species pool (Pärtel et al. 2013). Results suggest for those regions 308 where biodiversity in favourable species increased, land-use management may be more in-line 309 with biodiversity conservation objectives, and therefore be regions of conservation importance. 310 Conversely, declines in favourable species, reported among regions where overall biodiversity 311 declined, highlights regions that may benefit from biodiversity restoration. 312 It is possible for positive changes to have been borne, in part by the adoption of

313 traditional management funded by the Environmentally Sensitive Areas (ESA) scheme. 314 Introduced to Scotland, in 1987, ESA schemes provided financial incentives to land managers 315 to undertake environmentally friendly practices to maintain and enhance biodiversity, 316 landscape and historic values of designated habitats. Although the schemes have since been 317 superseded, for several regions where they were implemented, species richness increased (Fig. 318 6). The pattern suggests that financially incentivised, low intensity management may have 319 either; directly or indirectly begun a process of widespread maintenance and enhancement of 320 the machair.

321 There is however an exception to this pattern as more northern ESA designated areas 322 (Benbecula, North Uist and Shetland), displayed significant declines in favourable species. 323 This raises debate over the appropriateness of preservation policies that specifically aim to 324 preserve traditional management by making them financially attractive. In the EU, such 325 policies are implemented through the Common Agricultural Policy (CAP, European 326 Commission for Agriculture and Rural Development 2010), and their application is widely 327 unquestioned. However, there are concerns that such artificial ties between people and the 328 landscape will only lead to a decoupling between ecosystems and social systems (Fischer et al. 329 2012). Over time, naturally beneficial feedbacks between nature management and society are 330 forgotten, and where monetary incentives cease so do the environmentally beneficial land-use 331 practices. Biodiversity declines reported in this study for regions once designated as ESAs may 332 then have rapidly reverted towards more economic, yet environmentally insensitive land-use 333 practices.

334 Biodiversity gains alone may not necessarily be enough to conclude current land-use 335 trends to be inline with biodiversity conservation objectives. Many regions where favourable 336 species increased, species not belonging to the machair species pool (i.e. non-favourable 337 species) increased. This pattern was significant for three regions (Coll, Islay and Tiree), for 338 which species indicative of poor quality machair and active cultivation also increased. Results 339 suggest, for these regions arable cultivation may have intensified supporting the finding of 340 Lewis et al. (2013), where declines in functional richness were subsequently linked to land-use 341 intensification.

Changes in the frequency and or intensity of disturbances can result in a relaxation in the competitive interactions among plant communities, permitting successful invasions of species from the regional species pool (Mayfield et al. 2010). Where disturbances are intensified, invaders commonly possess ruderality or opportunistic life strategy traits, and often do not belong to the invading habitat species pool. Equally, many extant species, particularly

347 long-lived pants will possess high inertia to environmental change, capable of persisting in a 348 community as remnant populations long after changes have occurred (Diamond 1972; Eriksson 349 1996; Helm et al. 2006; Lindborg 2007). Therefore, at the regional-scale, invasions may not 350 result in an immediate loss of favourable species from the habitat-specific species pool. The 351 result is a colonization credit persisting of non-favourable species and a potential extinction 352 debt (i.e. extant species expected to go extinct because threshold conditions for their persistence are no longer satisfied; Tilman et al. 1994) persisting of favourable species. 353 354 Extinction debt has significant implications for conservation and restoration of threatened 355 grassland habitats and species (Helm et al. 2006; Cousins 2009). Machair regions where 356 biodiversity declines are not evident, but do show evidence of increased non-favourable 357 species linked to land-use intensification should therefore be of high restoration importance, 358 and warrant further investigations before desired parts of diversity are lost.

359

#### 360 **4.2. Compositional change: Linking shifts to environmental surrogates**

For grassland communities, functional characteristics of plant species can help to 361 362 distinguish between different land-use disturbances (Garnier et al. 2007, Pakeman 2011) and 363 land-use changes (Lavorel and Garnier 2002; Lewis et al. 2013). In this study we use very 364 simple functional classifications that differentiate between competitors, ruderals, stress 365 tolerators (Grime 1974), as well as heliophyte, helophyte, halophyte, nitrophilous, ericaceous 366 and calcicole species (Ellenberg 1988). Projection of these environmental surrogates on the 367 NMDS ordination, coupled with the simple proxy measures of disturbance, provides an 368 indication of land-use drivers of plant compositional change. We found significant national and 369 regional compositional changes, measured as shifts in Bray Curtis dissimilarity (Bray and 370 Curtis 1957) between survey years, suggesting a change in species community dynamics over 371 time. Positive changes along NMDS Axis 1 scores, for which was the most common 372 significant pattern among regions, could be linked to gains in ruderal species. As mentioned

373 earlier, invasion of ruderals is often a result of a decline in competitive interactions, usually 374 attributable to some form of disturbance. Three regions (Coll, Islay and Tiree), where positive 375 shifts along NMDS Axis 1 were identified, are also those regions identified in the biodiversity-376 change analyses to have likely suffered from agricultural intensification since the original 377 survey. Although not clear from the NMDS ordination, regression of change in land-use 378 proxies with change in ordination scores confirm arable cultivation practices to significantly 379 drive compositional change along NMDS Axis 1, and therefore subsequently linked to 380 significant compositional turnover of Scottish machair grasslands.

381 Correlates with NMDS Axis 2 were shown to be environmental surrogates linked to 382 proxies of disturbances. Livestock grazing was negatively ranked, while cultivation and 383 fertilisation practices were positively ranked. Land-use intensification through agricultural 384 intensification, particularly in conjunction with nitrogen deposition typically results in 385 compositional and taxonomic turnover among species assemblages, as environmental 386 conditions change to favour more competitive, fast growing species that locally displace slow 387 growing stress tolerators (Ordonez et al. 2009; Maskel et al. 2010). This pattern was evident 388 for one region (Shetland) that displayed a significant positive shift along NMDS Axis 2, a 389 significant decrease in stress tolerator species in favour of competitors and a significant gain in 390 Ellenberg N, indicative of an increase in fertility. The national increase in Ellenberg N, is 391 consistent with other long-term studies which report nation-wide gains in fertility values 392 among semi-natural vegetation types across much of the UK (McCollin et al 2000; Smart et al. 393 2005). Experimental studies on calcareous grasslands have found N deposition to lead to shifts 394 in species composition, with increases in rank grasses and loss of forbs and bryophytes 395 (Morecroft et al., 1994; Lee and Caporn, 1998; Johnson et al. 1999). However, regional gains 396 in Ellenberg N in this study were often shown with gains in competitive species without 397 biodiversity loss. It may be regions are experiencing a relaxation period between land-use changes and stable equilibriums (Diamond 1972), whereby local extinctions are likely for those 398

399 species with lower soil nitrogen preferences (Sarr et al 2012). However, it may equally be due 400 to interacting effects of multiple land-use practices, arable agriculture and livestock grazing, 401 whereby grazers prevent potential dominants taking advantage of the increases in fertility. 402 Increased fertility among semi-natural grasslands therefore present concerns for 403 biodiversity conservation, particularly where land-use disturbances through traditional farming 404 practices are in decline. Declines in active management, particularly in grazing intensity, 405 typically result in increased dominance of competitor species and decline in ruderal species as 406 competition for space, light and nutrients increase (Pakeman 2004). For many European 407 countries, cessation in active management is recognised as the major threat to semi-natural 408 grasslands (Levin 2012). Evidence of management declines were certainly prominent in our 409 study, illustrating where shifts in functional indicators suggest reduced disturbance, these 410 regions were mostly those with significant declines in biodiversity. This highlights the negative 411 impact declines in active management and disturbance intensity have upon the conservation 412 value of semi-natural grasslands. For the machairs of Scotland, it appears a complete de-413 coupling between the land and land-users to have a more immediate impact on biodiversity 414 value, than land-use intensification, stressing the importance of maintaining strong linkages 415 between nature and societies. Therefore, among landscapes with historic land-use traditions a 416 vital goal of conservation policy should be to help reconnect people with nature, fostering 417 socio-economic development while maintaining biodiversity (Fischer et al. 2012). 418

## 419 **4.3.** Value of biotic indicators and environmental surrogates for biodiversity monitoring

Given high temporal variability in land-use changes, and the ability of many species to persist long after disturbances have occurred, biodiversity indices alones reveal little information on whether current land-uses are in-line with conservation efforts. Interpretation of results becomes more spurious where sample efforts are small and temporal lags are large. However, introducing simple environmental surrogates as an additional component to

425 traditional diversity indices and select target species, help significantly for attributing land-use 426 drivers to observed patterns of change. Although rare, good quality temporal datasets can 427 provide documented evidence of vegetation composition prior to major global change impacts. 428 and are consistently proven for quantifying long-term patterns of vegetation change (McCollin 429 et al. 2000; Smart et al. 2005; Bennie et al. 2006). Given the extensive sampling effort in this 430 study non-representative samples should not have undue influence, however we do 431 acknowledge that caution should be expressed in the interpretation from only two snapshots in 432 time, particularly with large temporal lags between surveys.

This study clearly demonstrates the value of simple biotic indicators and environmental 433 434 surrogates for use in biodiversity monitoring assessments, demonstrated here across a large 435 national extent of semi-natural grasslands. In this study we used very simple indices and 436 functional indicators, which allowed us to differentiate between possible regions and infer with 437 some confidence the likely land-use changes driving species compositional and biodiversity 438 patterns. Our approach shows that functional indicators need not be overly complex, 439 particularly for assessing broad-scale regional change. Functional traits measures that are 440 nowadays also easily accessible are exceptional useful for gaining in depth knowledge of land-441 use impacts on the composition and biodiversity of plant assemblages, and can also be used to 442 identify community assemble processes (refs). Nevertheless, high variability in terms of land-443 use practices, land-use change and species compositions within single regions complicate the 444 generalisations that can be made. Therefore, we argue that broad functional classifiers to be 445 equally proficient in identifying patterns and process of change at broader spatial scales, and 446 are a useful tool for identifying regions of conservation concern that may warrant more 447 detailed, finer-scaled investigations.

Scale therefore is an extremely important factor when making interpretations from
biodiversity monitoring studies. Many pattern and trends identified at regional-scales were not
apparent at the national scales, as contrasting divergent and convergent trends between

451 geographic regions counteract, revealing little or no national pattern. Regional results from this 452 study are still generalisations, and are unlikely to reflect entire grassland assemblages within a 453 predefined geographic boundary. Nevertheless, our results clearly show the value to analysing 454 national data at smaller geographic scales.

455

## 456 5. Conclusions

457 This study clearly demonstrates the value of simple biotic indicators and environmental 458 surrogates for use in biodiversity monitoring assessments, demonstrated here across a large 459 national extent of semi-natural grasslands. We used very simple indices and functional 460 indicators, which allowed us to differentiate between possible regions and infer with some 461 confidence the likely land-use changes driving species compositional and biodiversity patterns 462 of Scottish semi-natural coastal grasslands. Our approach highlights that functional indicators 463 need not be overly complex, particularly for assessing broad-scale regional change. Functional 464 traits measures that are nowadays relatively easily to obtain, are also exceptionally useful for 465 providing an in-depth understanding of land-use impacts on the composition of vegetation 466 (Pakeman et al., 2009; Lewis et al., 2014), as well as insights into the community assembly 467 processes that govern plant assemblages (e.g. Mouchet et al., 2010). Nevertheless, high 468 variability in terms of land-use practices, land-use change and species compositions within 469 single regions complicate the generalisations that can be made. Simple, broad functional 470 classifiers, as used here, can therefore be equally proficient in identifying patterns and process 471 of change at broader spatial scales, and are a useful tool for identifying areas of conservation 472 concern that may warrant more detailed, finer-scaled investigations.

473

474 Spatial extent is therefore an extremely important factor when making interpretations 475 from biodiversity monitoring studies. In this study, many patterns and trends identified at the 476 regional scale were not apparent at the larger national scale, as contrasting divergent and 477 convergent trends between geographic regions counteracted, resulting in little or no national 478 pattern. Regional results from this study are still generalisations, and are unlikely to reflect 479 entire grassland assemblages within a pre-defined geographic boundary. Nevertheless, our 480 results clearly showed the value of analysing

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486

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Disturbance Proxy	Data description	Environmental Variable	Abbreviation
Animal signs	Plot level presence or	Cattle	N/A
C	absence of dung from	Sheep	N/A
	5 herbivore species	Rabbits	N/A
Grazing pressure	Intensity of herbivore	None	NoGrz
	grazing pressure	Light	LghtGrz
		Moderate	ModGrz
		Heavy	HvyGrz
Vegetation structure	Indication of vegetation height and structure for each plot $(0,1,2,3 = no,$ sparse, intermediate and full cover respectively)	Grasses < 20cm	Gra20
		Grasses 20-50cm	Gra2050
		Grasses > 50cnm	Gra50
		Herbs < 20cm	Hrb20
		Herbs 20-50cm	Hrb2050
		Herbs > 50cnm	Hrb50
Cultivation	Plot level presence or absence of a suite of cultivation factors,	No Cultivation	NoCult
		Standing Crop	StdCrp
		Recent Cultivation	RecCult
	from no cultivation	(past 3 years)	
	to current standing crop	Old Cultivation (4 or more years)	OldCult

**Table 1:** Description of land-use proxies measured for each 5m x 5m plot.

685

**Table 2.** National and regional Mean ( $\overline{X}$ ) temporal shift in the species diversity of machair grassland between 1976 and 2010 measured through i. richness (species count) and ii. Inverse Simpson index. Significance tested through paired t tests (national) and general liner mixed model with poisson function (regional). '\*\*\*'  $p \le 0.001$ , '\*\*'  $p \le 0.01$ , '\*'  $p \le 0.05$ .

					691		
	Speci	Species Richness			1/Simpsons		
	$\overline{x}$	Z	sig	$\overline{x}$	z 69122		
National	2.32	7.60	***	-0.25	2.17 *		
Barra	4.21	4.31	***	-2.33	-4.28 <b>693</b> *		
Benbecula	-3.36	-1.99	*	-1.71	-1.97 *		
Coll	10.07	12.51	***	0.67	1.72 694		
Colonsay	14.70	7.39	***	2.68	2.75 **		
Harris	-5.13	-3.60	***	-2.44	-3.12 695		
Islay	8.89	8.47	***	0.22	0.40		
Lewis	-2.15	-2.59	**	0.40	1.02 696		
Monach	-3.06	-2.87	**	-1.46	-2.97 **		
North Coast	-1.02	-1.33		1.15	3.44 6 <b>9</b> 7*		
North Uist	-0.35	-0.68		-1.30	-5.16 ***		
Sanday	0.04	0.04		0.17	0.38 698		
Shetland	-3.29	-2.97	**	-1.08	-1.89		
South Uist	4.98	7.40	***	-0.93	-2.85 699		
Tiree	10.23	13.96	***	1.96	5.34 -***		
					/00		

**Table 3.** Significance of change in land-use disturbance proxies in explaining temporal compositional change as measured by the NMDS analyses (Figure 3). Significance tested through general liner mixed model. '\*\*\*'  $p \le 0.001$ , '\*\*'  $p \le 0.01$ , '\*'  $p \le 0.05$ .

						705
	NMI	DS AXIS	51	NMI	OS AXIS	$52^{00}$
Land-Use						706
NoGrz	0.05	0.66		0.02	0.42	
LightGrz	0.05	0.76		0.03	0.77	707
ModGrz	0.03	0.37		0.03	0.57	
HvyGrz	0.05	0.66		0.06	1.28	708
RecCult	0.48	5.69	***	0.28	4.97	***
OldCult	0.17	3.02	**	0.11	3.11	709
StndCrp	0.10	0.68		0.08	0.83	
Gra20	-0.02	-1.08		< 0.01	< 0.01	710
Gra2050	-0.01	-0.65		-0.01	-1.13	
Gra50	-0.06	-3.23	***	-0.01	-1.17	711
Hrb20	0.04	2.16	*	0.01	0.86	
Hrb2050	0.03	1.65		< 0.01	0.35	712
Hrb50	0.06	1.67		0.05	2.31	*
Cattle	< 0.01	0.03		0.02	1.07	713
Sheep	0.01	0.30		-0.02	-1.00	
Rabbit	0.01	0.38		-0.01	-0.50	714

Fig. 1. Distribution of original and resurveyed study sites across Scotland. For site details seeAppendix A.

718

719	Fig. 2. Mean richness and % cover change for species groups indicative of different conditions
720	and land-use states of machair grasslands: Cult = species indicative of cultivated land; Fallow
721	= species indicative of rotational arable machair i.e. fallow grasslands; Machair = species
722	indicative of machair grasslands; Poor = species indicative of degraded machair grasslands.
723	Mean differences are presented with standard error bars of the mean. Shaded bars represent
724	statistically significant change ( $p < 0.05$ ) tested using General Linear Mixed Models (GLMM;
725	Richness) and linear-mixed modelling with residual maximum likelihood (REML;
726	Abundance).
727	
728	Fig. 3. NMDS ordination of machair grassland communities across Scotland, computed using
729	the Bray Curtis dissimilarity. Figures i-ii: Displayed are the convex hulls for each survey
730	(dashed and solid lines refer to the original and resurvey respectively). (i) Projected
731	environmental proxies for land-use p<0.001 (see Table 1). (ii) Projected functional signatures:
732	Ellenberg and CSR scores. (iii) Displayed as text are species with a significant statistical
733	relationship (p < 0.001) and a goodness of fit ( $r^2 \ge 0.05$ ) along Axis 1 and or Axis 2. Species
734	are represented by the following letter codes: Aa Ammophila arenaria, As Agrostis stolonifera,
735	Bry Bryophytes, Cf Carex flacca, Cn Carex nigra, Cp Cardamine pratensis, Er Elymus repens,
736	Gp Galium palustre, Gv Galium verum, Hv, Hydrocotyle vulgaris, Lp Lolium perenne, Pa
737	Potentilla anserina, Pv Prunella vulgaris, Rr Ranunculus repens, Sp Succisa pratensis, Tp
738	Thymus polytrichus, Tr Trifolium repens. Non-significant species are represented by '•'.
739	
740	Fig. 4. National and regional shifts in NMDS ordination axes scores for computed using Bray

741 Curtis in two dimensional ordination space. Displayed are kernel densities for NMDS axis 1

742 and axis 2. Dashed and solid lines refer to the original and resurvey respectively. Shaded 743 density graphs (green = original survey & purple = resurvey) represent statistically significant 744 shifts between survey years (p < 0.05) tested using nested linear mixed models with residual 745 maximum likelihood (REML). 746 747 Fig. 5. National and regional cover-weighted functional classification values (i. CSR & ii. 748 Ellenberg) for the surveyed machair grasslands of Scotland. Mean plot scores are presented 749 with standard error bars of the mean. Grey bars represent statistically significant change 750 between survey years (p < 0.05) tested using nested linear mixed models with residual 751 maximum likelihood (REML).

752

Fig. 6. Patterns between regional species richness gains and loses since 1976 and those regionsonce designated as an Environmentally Sensitive Area (ESA).