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Effects of adding an arbuscular mycorrhizal fungi inoculum and of distance to donor sites on plant species recolonization following topsoil removal

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1 Effects of adding an arbuscular mycorrhizal fungi inoculum and of distance

2 to donor sites on plant species recolonization following topsoil removal

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Questions: Does addition of an arbuscular mycorrhizal fungi (AMF) inoculum increase the short-term restoration success of a nutrient-poor grassland (NPG) after topsoil removal? Does distance to intact remnant grassland (IRG) patches affect the restoration success, and does the effect of inoculum addition depend on the distance to IRGs?

22 **Location:** Meerdaal forest, Oud-Heverlee, Belgium.

23 Methods: In a topsoil-removed site of 8.5ha, where 24 IRG patches (c. 10% of the area) were kept, 48 plots (1m²) were established at three distances (5, 10 24 25 and 20m) from the edge of IRG patches. Half of the plots at each distance class 26 were inoculated with a custom-made AMF-inoculum, whereas the remaining 27 were used as controls. We recorded the plant species abundance in the plots, 28 just before the addition of the AMF-inoculum, and one year after. We used 29 repeated measures ANOVAs to test for effects of inoculum addition, distance to 30 the IRG patches, and their interaction, on plant species richness, diversity, and 31 on the plant community similarity between IRG patches and plots. We also evaluated the response of AMF-dependent plant species, specialist plant species
 of NPG, and plant species with specific seed dispersal adaptations.

34 **Results:** Adding the inoculum positively affected the species richness and/or 35 diversity of all plant species, AMF-dependent plant species and specialist plant 36 species. It increased plant community similarity to the IRG patches. Increasing 37 distance from the IRG patches had a negative effect on the richness and/or 38 diversity of all plant species and specialist plant species. The positive effect of 39 inoculum addition on richness and/or diversity of all plant species, AMF-40 dependent plant species and specialist plant species decreased with increasing 41 distance from the IRG patches to the plots, likely indicating priority effects.

42 **Conclusions:** The application of a custom-made AMF-inoculum increased the 43 short-term restoration success of NPG after topsoil removal. Dispersal limitation 44 of specialist plant species of NPG, however, likely negatively affected the effect 45 of inoculum addition. Apart from the AMF-effect, the reported strongly positive 46 short-time effect of the inoculation was likely due to the high density of IRG 47 patches at the site, and to the presence of organic-matter and other micro-48 organisms in the inoculum.

49

Keywords: AMF inoculum; dispersal limitation; dispersal syndromes; ecological
 restoration; nutrient-poor grasslands; priority effects.

52

53 **Nomenclature:** van der Meijden (2005).

54

55 Introduction

56 Owing to a widespread decline of species-rich semi-natural grasslands in Europe 57 (Poschlod & Wallis De Vries 2002), ecological restoration of these habitats has 58 been a priority since the 1970ies (Bakker 1989; Walker et al. 2004). These 59 activities have also been strongly supported by the European Habitat directive 60 (92/43/EEC), which aims at protecting and conserving habitats and wild fauna 61 and flora. By restoring degraded ecosystems to their previous state, both species 62 diversity and ecosystem functioning can recover as well (Brudvig 2011). Whether or not successful restoration is achieved, largely depends on a range of biotic 63 64 and abiotic constraints (Walker et al. 2004; Cramer et al. 2008), especially regarding soil properties and the capacity of target plant species to recolonize. In 65 66 this respect, it has been shown that high soil nutrient levels constrain the 67 successful restoration of nutrient-poor ecosystems (Fagan et al. 2008, 2010), 68 whereas dispersal limitation (Standish et al. 2007; Öster et al. 2009) and the 69 absence of soil biota (Kardol et al. 2006; Vergeer et al. 2006) may limit the arrival 70 and establishment of seedlings of the target species (Öster et al. 2009).

71 Soil nutrient enrichment is largely caused by anthropogenic activities such 72 as fertilization and atmospheric deposition from combustion processes (Peñuelas 73 et al. 2012), and has been shown to strongly affect the occurrence and 74 community composition of plant species. For instance, several studies have 75 reported lowered species richness and shifts to dominance of a few competitive 76 species in grasslands under increased nitrogen (N) input (Bobbink et al. 2010) 77 and under phosphorus (P) enrichment (Ceulemans et al. 2013, 2014). In order to 78 reduce nutrient levels at ecological restoration sites, high intensity interventions 79 such as topsoil removal are often the only solution, especially for removing the 80 immobile soil-P (Weijtmans et al. 2009; Pedley et al. 2013). However, Geissen et 81 al. (2013) has recommended against using topsoil removal as a nature 82 management technique due to its negative effects on soil quality and soil biota. It 83 has indeed been reported that after topsoil removal only a small subset of the 84 target plant species was able to recolonize the restoration sites (Verhagen 2007; 85 Bekker 2008), even when target plant species were occurring in adjacent communities (Dobson et al. 1997; De Graaf et al. 1998; Bakker & Berendse 86 87 1999; Vergeer et al. 2006). Failure of target plant species to establish, despite 88 low seed dispersal constraints may result from a lack of soil biota such as 89 arbuscular mycorrhizal fungi (AMF) (van der Heijden 2004; Verhagen 2007), 90 which are generally removed with the topsoil (Vergeer et al. 2006).

91 Arbuscular mycorrhizal fungi form a symbiosis with >80% of the land plant 92 species (Smith & Read 2008). The symbiosis is based on a reciprocal exchange

93 of resources: the host plant provides photosynthates to the fungus, and in return 94 it receives vital inorganic nutrients (Smith & Read 2008). AMF may also 95 contribute to seedling establishment by integrating the seedlings into an existing 96 mycorrhizal network connected with already established adult individuals (Simard 97 & Durall 2004; van der Heijden & Horton 2009). Furthermore, the symbiosis 98 induces metabolic changes in the hosts through impact on defense hormone 99 production (Pozo & Azcón-Aguilar 2007; Jung et al. 2012). As a consequence, 100 plant resistance against soil pathogens (Whipps 2004), nematodes (De la Peña 101 et al. 2006), and abiotic stress, such as drought, salinity and heavy metals (Miransari 2010; Smith et al. 2010) is increased. 102

103 Clearly, AMF are vital to plant communities. Yet once absent, 104 spontaneous re-colonization of AMF occurs very slowly (Allen and Allen 1992), 105 as dispersal happens through colonization of roots from plant to plant (Read et 106 al. 1976) and through transport of spores by wind (Egan et al. 2014), small 107 mammals (Fracchia et al. 2011), water (Walker 1988), or soil fauna (Klironomos 108 & Moutiglis 1999). The natural restoration of the entire fungal community may 109 take decades (Baar et al. 2008). Vergeer et al. (2006) reported that two and a 110 half years after topsoil removal, the abundance of AMF spores corresponded to 111 55-70% of the AMF spore numbers found in natural grasslands. Addition of AMF 112 inoculum may therefore be crucial to accelerate the development of target plant 113 communities at restoration sites that have been topsoil stripped.

114 In addition to local biotic and abiotic conditions, plant assembly also 115 strongly depends on the landscape context of the restoration site. Evidence 116 shows differential effects of distance to seed sources on plant species 117 recolonization, including strong effects (Bischoff et al. 2009; Pottier et al. 2009), 118 and weak or non-existing effects (Cole et al. 2010; Matthews & Endress 2010). 119 Furthermore, Helsen et al. (2013) found that spatial isolation filters plant species 120 based on their dispersal capacity, slowing down the community assembly 121 process towards the target community. Priority effects occur when earlier arriving 122 species affect the establishment, growth or reproduction of later arriving species 123 (Harper 1961; Chase 2003). These effects can lead to long-lasting differences in species composition between the target community and the restored community.
Therefore, the spatial context of the restoration site should be incorporated into
the restoration design (Matthews et al. 2009).

127 The aim of this study was to test the effectiveness of the application of a 128 custom-made AMF inoculum in increasing the short-term restoration success of 129 nutrient-poor grasslands where the topsoil had been removed. We hypothesized 130 that the addition of AMF inoculum would enhance establishment success of 131 AMF-dependent plant species. Additionally, we aimed at testing how the spatial 132 context of the restoration site, in relation to intact remnant grassland (IRG) 133 communities that can act as a plant species source, affect the restoration 134 success. We tested the hypothesis that the establishment success of specialist 135 plant species of nutrient-poor grasslands, and of plant species with poor seed 136 dispersal capacities, would decrease with distance from the IRG. Finally, we 137 evaluated whether the effect of the addition of the AMF inoculum depended on 138 the distance to the IRG patches. Here, we hypothesized that colonization by 139 AMF-dependent plant species will decrease with increasing distance from the 140 IRG patches, because well-dispersed plant species will monopolize distant 141 restoration sites, thus negatively affecting the effectiveness of the AMF inoculum 142 in distant plots.

143

144 Methods

145 Site description

146 The study was carried out in a large clearing in Meerdaal forest in central 147 Belgium, south of Leuven (50° 48' 32.15" N, 4° 40' 34.42" E, 78 m a.s.l.). The 148 study site was a former military domain (c. 8.5 ha) with ammunition storage 149 facilities. In the context of the European habitat directive (92/43/EEC), a nature 150 conservation project was carried out to restore nutrient-poor grasslands and 151 grassy heathlands. After the removal of the remaining military buildings and 152 roads, the topsoil was removed in January 2011, to a depth of 10 cm, to remove 153 the nutrient-rich topsoil layer, down to the mineral soil. At the restoration site, 24 154 scattered patches of well-developed grassland communities were present around the former ammunition storage locations. These IRG patches (sized between 0.03 and 0.04 ha, in total c. 10% of the area) occur in a regular grid pattern, and contain the original species-rich vegetation with characteristic and regionally endangered species of nutrient-poor grasslands such as *Polygala serpyllifolia*, *Thymus pulegioides* and *Campanula rotundifolia*.

160

161 Experimental design

162 In mid-August 2012, a total of 48 permanent plots of 1 m x 1 m were randomly 163 established at three distances (16 replicates at each distance of 5, 10 and 20 m) 164 from the edge to the closest IRG patch. Eight out of the 16 plots at each distance 165 class were inoculated with a custom-made AMF inoculum (see further), whereas 166 the remaining eight plots were not inoculated and served as controls (Fig. 1). To 167 apply the inoculum, we removed the upper 1 cm layer of soil with a hand rake 168 without affecting the sparse already established plants in the plot. We applied 169 900 ml of the custom-made liquid AMF inoculum per plot. After the application of 170 the inoculum, we covered the plots with the removed soil. The plots that did not 171 receive the inoculum addition were submitted to the same soil disturbance with 172 the hand rake and de-mineralized water was applied.

173

174 **Preparation and ex situ testing of the AMF inoculum**

175 AMF-colonized roots can be used as a source of AMF inoculum (Tommerup 176 1984; Klironomos & Hart 2002). Because the origin of the AMF may strongly 177 influence their effectiveness (White et al. 2008; Pellegrino et al. 2011), we used 178 the roots of AMF-dependent plant species collected in October 2011 in the wide 179 surroundings of the study area. These plant species are indicator species of 180 nutrient-poor grasslands that are still relatively common, and included Achillea 181 millefolium, Centaurea jacea, Danthonia decumbens, Hieracium pilosella, 182 Hypochaeris radicata, Potentilla erecta, Stachys officinalis, and Succisa 183 pratensis. Roots were soaked and washed out of soil with tap water, but still 184 some soil remained attached. Afterwards, roots were blended separately per 185 species in de-mineralized water with a commercial Philips HR7625 food processor to chop them in small pieces, between 0.1 and 3 cm. The resulting
muddy mixture of each species was equally distributed across 24 containers of
300 ml.

189 A greenhouse experiment was then ran to test the effectiveness of this 190 custom-made inoculum through assessments of AMF root colonization and spore 191 density, and to test if it contained pathogens and seeds that may confound the 192 spontaneous colonization process. The experiment ran during four months. 193 Three known AMF-dependent plant species (Campanula rotundifolia, Nardus 194 stricta, and Festuca filiformis) were seeded as monocultures in 2 L pots, each pot 195 containing 15 seeds. We used a substrate composed by potting soil and sand 196 (3:1, respectively), both were autoclaved at 120 °C for one hour. The plant 197 species were sown on sterilized substrate (control) and on sterilized substrate 198 inoculated with the inoculum. Each treatment was replicated eight times, 199 resulting in a total of 48 pots. We applied 300 ml of the AMF inoculum to the pots 200 and de-mineralized water to the control pots. Root colonization and spore density 201 were surveyed to test the effectiveness of the inoculum. After four months, roots 202 were washed and stained following Grace & Stribley (1991), with 1:1 1% Methyl 203 blue solution and 85% Lactic acid. Root samples were observed under a 204 dissecting microscope at 32x magnification. The grid line intersect method 205 (Giovannetti & Mosse 1980) was used to measure the degree of AMF 206 colonization. In addition, 100 g of soil per pot was collected to extract the AMF 207 spores, following the sucrose centrifugation method of Brundrett et al. (1994). 208 Spores were counted using a dissecting microscope at 32x magnification. We 209 found that the roots of all individuals that were inoculated were colonized by 210 AMF. The mean percentage of colonization was 57.4% for C. rotundifolia, 50.4% 211 for N. stricta and 25.1% for F. filiformis. In the inoculated pots, we counted between 49-207 spores g⁻¹ soil; none of the plants developed any type of 212 213 disease. There were no AMF spores, and there was no AMF root colonization in 214 the non-inoculated control pots. In addition, only two pots out of the 24 inoculated 215 pots had non-sown Juncus effusus growing, which demonstrates a very low 216 probability of seed introduction with the inoculum. In July 2012, the same procedure was then followed to prepare the AMF inoculum to be used at thestudy site.

219

220 Vegetation surveys

221 In early August 2012, just before adding the AMF inoculum, we recorded the 222 plant species abundance in the established 1 m x 1 m plots. A grid, consisting of 223 100 10 cm x 10 cm squares was placed on top of the plots and we counted the 224 number of individuals of plant species that occurred in each square. The same 225 procedure was repeated in July 2013. Additionally, we recorded all plant species 226 occurring in each of the 24 IGR patches (Table S1. Supplementary material). In 227 both surveys, 18% of the individuals were identified at family or genus level due 228 to their early growth stage.

229

230 Data analyses

231 For both vegetation surveys and for each plot, we calculated both plant species 232 richness and the Shannon diversity index to the power of e (Jost 2006) for all 233 plant species, for AMF-dependent plant species (following Fitter & Peat (1994)), 234 and for specialist plant species of nutrient-poor grasslands (following Decleer 235 2007; see Table S1, Supplementary material). In addition, we calculated the 236 plant community similarity between the IRG patches and the experimental plots, 237 using the Jaccard similarity index. Non-metric multidimensional scaling (NMDS) 238 was used to analyze the similarity in plant community composition among plots. 239 The NMDS analysis was based on the plant species abundance x plot matrix of 240 the two surveyed years together, and on Bray-Curtis dissimilarities, with several 241 starting points and a maximum of 1000 iterations. These analyses were done 242 separately for (i) all plant species, (ii) AMF-dependent plant species, and (iii) 243 specialist plant species. Furthermore, we determined the plant species richness 244 according to plant dispersal syndromes. We distinguished between 245 endozoochores, myrmecochores and anemochores (Kleyer et al. 2008; hereafter 246 referred to as plant species dispersed by dung, ants and wind, respectively). The 247 plant species were assigned to the prevailing dispersal syndrome. The nine most widely distributed plant species in the plots were assigned to the category of welldispersed plant species. Plant species richness, Jaccard similarity index, and
NMDS were calculated using the R program 3.0.2 (R Foundation for Statistical
Computing, Vienna, AT), with the package *Vegan* (Oksanen et al. 2011).

Prior to further analyses, F-tests were performed on the vegetation data of the initial survey (2012) to confirm that there were no *a priori* differences between plots at different distances from the IRG patches. To meet assumptions of normality, several response variables were log or square root-transformed.

256 We then conducted repeated measures ANOVAs to test for effects of 257 inoculum addition (inoculated vs. non-inoculated), distance to the closest IRG 258 patch (5, 10 or 20 m), and their interaction, on the differences in plant species 259 richness, Shannon diversity, and Jaccard index between the surveys in 2012 and 260 2013. ANOVAs were run using the R package *nmle* (Pinheiro et al. 2009). 261 Backwards model selection was used. When a significant effect was found, 262 multiple comparisons were performed with the R package phia (De Rosario-263 Martinez 2012). Repeated measures ANOVAs were done separately for all plant 264 species, AMF-dependent plant species, and specialist plant species. Similar 265 analyses were done using species richness of the three plant dispersal syndrome 266 categories and the well-dispersed plant species, as the dependent variables.

Finally, the NMDS scores (on the first two axes) of the plots in 2012 and 268 2013 were used as measures of plant community similarity among plots. These 269 values were used as dependent variables in a repeated measures ANOVA, as 270 described above. Again, these analyses were done using the NMDS scores of all 271 plant species, AMF-dependent plant species, and specialist plant species.

272

273 Results

In total, we recorded 80 plant species in the IRG patches and 78 plant species in the plots, belonging to 28 families; 52 of them were AMF-dependent plant species, and 11 were nutrient-poor grasslands specialist plant species (Table S1). In 2012, we recorded 55 plant species; 37 were AMF-dependent plant species, and 9 were specialist plant species. In 2013, we registered 66 plant

279 species; 43 were AMF-dependent plant species, and 11 were specialist plant 280 species. Of the nine well-dispersed plant species, five were AMF-dependent 281 (Holcus lanatus, Hypericum perforatum, Juncus effusus, Molinia caerulea, Rubus 282 fruticosus), and four were not (Calluna vulgaris, Carex pilulifera, Luzula 283 multiflora, Pinus sylvestris). In 2012, the number of plant species recorded per 284 plot ranged from 1 to 28 (mean = 12, SD = 6.4), and in 2013 it ranged from 4 to 285 32 (mean = 15, SD=7). Prior to inoculation, no significant differences were found 286 among the three distance classes for plant species richness ($F_{2.45}$ = 2.6, P > 287 0.05), Shannon diversity ($F_{2.45}$ = 2.96, P > 0.05), Jaccard index ($F_{2.45}$ = 5.2, P > 288 0.05), NMDS1 ($F_{2.45}$ = 4.2, P > 0.05), and NMDS2 ($F_{2.45}$ = 4.5, P > 0.05).

289

290 Plant species response to treatments

291 Inoculation significantly increased total plant species richness, the Jaccard 292 species similarity index between the plots and IRG patches, and the plant 293 community similarities among inoculated plots (Table 1, 2). The effect of the 294 distance treatment on Shannon diversity was significantly higher in plots at 5 m 295 than in plots at 20 m (Table 1, 2). The effect of inoculum addition on plant 296 species richness, Jaccard similarity index, and plant community similarities 297 among plots varied with distance to the IRG patch (Table 1). These variables 298 were significantly higher in inoculated plots at 5 and 10 m than in inoculated plots 299 at 20 m, whereas no significant differences were found between inoculated plots 300 at 5 and 10 m (Fig. 2A, B, C). No significant differences were found among the 301 three distances in the non-inoculated plots for these response variables. Plant 302 species richness, Jaccard index between plots and IRG patches, and plant 303 community similarities among plots were significantly higher in inoculated plots at 304 5 and 10 m, as compared to non-inoculated plots at the same distances. 305 However, there was no difference among inoculated and non-inoculated plots at 306 20 m.

307 For the AMF-dependent plant species, inoculation significantly increased 308 richness, Shannon diversity and Jaccard species similarity with IRG patches 309 (Table 1, 2). The effect of the distance treatment was significant for community

similarities among all plots (Table 1). These similarities were higher in plots at 5 310 311 and 10 m than in plots at 20 m (Table 2). The effect of inoculum addition on plant 312 species richness, Shannon diversity, and the Jaccard index varied with distance 313 to the IRG patch (Table 1). Richness was higher in inoculated plots at 5 m than in 314 inoculated plots at 20 m (Fig. 3A), whereas no significant differences were found 315 among inoculated plots at 10 and 20 m, and among 5 and 10 m. Richness in 316 non-inoculated plots did not differ significantly among the distance classes. 317 Shannon diversity and the Jaccard index was significantly lower in inoculated 318 plots at 20 m, as compared to plots at 5 and 10 m, whereas no significant 319 differences were found between inoculated plots at 5 and 10 m (Table 2, Fig. 3B, 320 C). These response variables in non-inoculated plots did not significantly vary 321 with distance. Richness, Shannon diversity and the Jaccard similarity index with 322 IRG patches were significantly higher in inoculated plots at 5 and 10 m, as 323 compared to non-inoculated plots at same distances. However, inoculated and 324 non-inoculated plots at 20 m did not significantly differ.

325 For the specialist plant species of nutrient-poor grasslands, the inoculum 326 addition significantly increased the Shannon diversity and Jaccard similarity 327 index (Table 1, 2). The effect of the distance treatment on specialist plant species 328 richness and community similarities of specialist plant species among all plots 329 was significant (Table 1), both variables had higher values in plots at 5 and 10 m 330 than in plots at 20 m (Table 2). The effects of the inoculum addition on the 331 Jaccard index varied with distance from the IRG patches (Table 1). The Jaccard 332 index was significantly higher in inoculated plots at 5 or 10 m than in plots at 20 333 m (Fig. 4), while no significant difference was found between plots at 5 and 10 m. 334 The Jaccard index between non-inoculated plots and IRG patches was not 335 significantly different among the different distance classes. The Jaccard index 336 was significantly higher in inoculated plots at 5 and 10 m, as compared to non-337 inoculated plots at same distances. However, these similarities were not 338 significantly different in inoculated vs. non-inoculated plots at 20 m.

The inoculum addition significantly increased the richness of plant species dispersed by wind, ants or dung (Table 1, 2). The effect of inoculum addition

341 significantly varied with the distance treatment (Table 1); plant species richness 342 of the three groups was significantly higher in inoculated plots at 5 and 10 m than 343 in inoculated plots at 20 m (Fig. 5A, B, C). In non-inoculated plots the plant 344 species richness did not vary significantly among the different distance classes. 345 Plant species richness of these three groups was significantly higher in 346 inoculated plots at 5 and 10 m, as compared to non-inoculated plots at same 347 distances. However, plant species richness in inoculated vs. non-inoculated plots 348 at 20 m was not significantly different.

Finally, the effects of inoculum addition, distance and their interaction on NMDS2 scores were not significant for any of the studied groups of plant species (results not shown). Similarly, there were no effects of the two treatments and their interaction on the abundance of the well-dispersed plant species (results not shown).

354

355 **Discussion**

356 Effects of AMF-inoculum addition

357 The aim of this study was to test whether the application of an AMF inoculum 358 could improve the short-term restoration success of a nutrient-poor grassland 359 where the topsoil was removed. We predicted that the addition of the inoculum 360 would enhance the establishment success of AMF-dependent plant species. Our 361 results demonstrated that one year after topsoil removal, the plots that were 362 inoculated showed a higher Shannon diversity and richness of AMF-dependent 363 plant species, at least in the plots closest to the IRG patches that could act as 364 plant species sources. Zhang et al. (2012) found similar results, one year after 365 inoculating degraded grasslands in China with a lab propagated AMF inoculum.

The type of inoculum that we used is less expensive than commercial inoculum, and it is less time consuming to produce than inoculum that is produced by isolation of AMF spores and further propagation in plant roots. In addition, it is likely unfeasible to generate a pure inoculum consisting of many different and naturally occurring AMF taxa, because many of these are impossible or very difficult to cultivate and propagate. Although we have no 372 information on the specific AMF taxa that were present in the inoculum, we can 373 assume that they were very similar to the ones naturally occurring in nutrient-374 poor grasslands in the study region. AMF species composition is indeed known 375 to affect plant diversity and community composition, especially when a majority of 376 the plants in the community are AMF-dependent (Vogelsang et al. 2006). In 377 addition, White et al. (2008) and Pellegrino et al. (2011) experimentally 378 demonstrated that the origin of the inoculated AMF taxa has an important effect 379 on plant performance. These authors reported that inoculum produced through 380 propagation of AMF species from the local ecosystem was more effective in 381 improving plant performance than a commercially available AMF inoculum. The 382 use of native AMF from roots of plant species from undisturbed grasslands may 383 therefore be a convenient alternative to a commercial AMF inoculum, and may 384 offer important ecological and economic advantages.

385 It is important to note that our results cannot be exclusively attributed to 386 AMF in the inoculum, but also to the presence of organic matter and other soil 387 micro-organisms. Soil amendments have been proven to be beneficial in 388 ecological restoration (de Deyn et al. 2003; Carbajo et al. 2011), and to play a 389 facilitating role in establishing the soil microbial-plant association. Organic matter 390 amendments have also been suggested to stimulate and improve plant 391 mycorrhization (Douds et al. 2006, Jaison et al. 2011). In degraded alpine areas 392 in Switzerland, Schmid et al. (2008) found that through applying an (commercial) 393 AMF inoculum only, 65% of the area recovered with vegetation, in comparison to 394 87% when commercial AMF inoculum was applied along with organic nutrients 395 and P-solubilizing Penicillium spp. Nevertheless, the strong root colonization and 396 high density of AMF spores in our greenhouse experiment support the role of 397 AMF on our results. It was also the explicit objective of this study to evaluate a 398 practical and feasible ecological restoration approach, through adding a custom-399 made inoculum. It is currently indeed not possible to generate a 'pure' AMF 400 inoculum consisting of many different, naturally occurring AMF species only, 401 because many of these are impossible or very difficult to cultivate and propagate.

Such a pure inoculum would never mimic the AMF composition of our custom-made inoculum.

404

405 **Effects of distance to the donor sites**

406 We also tested how the distance to IRG that can act as a plant species sources 407 affected the restoration success. We predicted that the establishment success of 408 nutrient-poor grassland specialist plant species would decrease with distance 409 from the IRG. Our results show that increasing distance from IRG patches had a 410 negative effect on specialist plant species richness. Such step-wise dispersal is 411 consistent with the findings that many grassland specialists are poor dispersers 412 (Martin & Wisley 2006; Cousins & Lindborg 2008; Helsen et al. 2013). We did, 413 however, not find a distance effect on the Jaccard similarity index between 414 specialist plant species in the IRG patches and the study plots. This might be due 415 to the relatively low number of specialist plant species in the area (12 species in 416 total, range in the plots between 0 and 6 species). We also found no effects of 417 distance on the total plant species richness. Similarly, Krauss et al. (2004) found 418 no effects of distance on generalist plant species richness colonization in 419 calcareous grasslands in Germany. In contrast, Öster et al. (2009) reported a 420 decline of generalist plant species richness with increasing distance from 421 species-rich semi-natural grasslands to former arable fields, and Diacon-Bolli et 422 al. (2013) found that the amount of captured diaspores decreased abruptly at 20 423 m, in a seed rain study in a calcareous grassland. Also the Jaccard similarity 424 index between the IRG patches and the plots did not change with distance in our 425 study, suggesting that generalist plant species disperse and establish well across 426 the studied distances.

We also predicted that establishment success of ant-dispersed plant species, but not of wind- and dung-dispersed plant species, would decrease with distance from the IRG patches. We found that distance between the IRG patches and the study plots had no effect on the richness of plant species dispersed by wind, ant or dung. These results are consistent with the findings of Öster et al. (2009), who reported that wind- and animal-dispersed seeds were able to

successfully disperse up to 10 m from grasslands into former arable fields. That
we did not find a negative effect of distance on ant-dispersed plant species is
likely due to the very limited occurrence of such species (range in the plots
between 2 and 3 species) at our study site.

437

438 Effects of AMF inoculum addition depend on the distance to the donor site

439 Finally, we evaluated whether the effect of the addition of AMF inoculum on plant 440 community composition depended on the distance of the plots to the IRG. We 441 hypothesized that AMF-dependent plant species colonization would decrease 442 with increasing distance from the IRG patches because well-dispersed plant 443 species would monopolize the distant plots, obscuring the effectiveness of the 444 inoculation. We indeed found that in the inoculated plots, the richness of AMF-445 dependent plant species and the Jaccard similarity index between the IRG 446 patches and plots was relatively constant up to a distance of 10 m, whereas it 447 decreased at 20 m. On the other hand, in the non-inoculated plots, no significant 448 differences were found along the studied distance classes. This pattern of 449 community similarities demonstrates that the positive effect of inoculum addition 450 is reduced in distant plots. We also found that the abundance of well-dispersed 451 plant species did not vary across the studied distances, additionally suggesting 452 that the effectiveness of the inoculation at 20 m distances is obscured by 453 dispersal limitation of AMF-dependent plant species, in combination with priority 454 effects exerted by well-dispersed generalist plant species. Interestingly, the latter 455 was composed by AMF-dependent and non AMF-dependent plant species, then 456 the priority effects exerted over the other AMF-dependent plant species were 457 independent to AMF-dependency. This suggests that well-dispersed generalist 458 plant species created soil legacies, and these legacies contributed to strong 459 priority effects on AMF-dependent plant species (Grman & Suding 2010).

460 Our findings suggest the high ecological significance of inoculation in early 461 stages of restoration. Even though the positive effect of the addition of AMF 462 inoculum decreases in distant areas due to dispersal limitation, the addition of 463 AMF inoculum should be a desirable part of ecological restoration and can be 464 considered as a best practice in ecological restoration, likely along with seed 465 addition. Further work should focus on elucidating AMF species composition of 466 roots of plants from inoculated and non-inoculated plots, for example using 467 amplicon-sequencing approaches (e.g. Van Geel et al. 2015), in order to 468 disentangle the role of AMF addition and the addition of organic matter and other 469 soil micro-organisms.

470

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Table 1. Inoculation strongly positively affects plant species colonization of725topsoil removed sites, but the effect of inoculation is generally highly dependent726on the distance from the donor site. *t*-values from repeated measures ANOVAs727evaluating the effects of inoculum addition, distance to remnant grassland728patches, and their interaction on responses of different plant species sets.729Significance: ***, P < 0.001; **, P < 0.01; *, P < 0.05.

Species set	Variables	Inoculum	Distance	Ino y Dist
	Variables	(Ino)	(Dist)	
	Species richness	2.7***		-3.3***
	Shannon		2 0**	
All plant	diversity		-2.0	
species	Jaccard index	4.8***		-5.7***
	NMDS1	3.2**		-3.6***
	'Species richness	2.5**		-2.8**
	Shannon	2.3*		-2.8**
AMF-	diversity	2.0		2.0
dependent plant species	Jaccard index	4.1***		-4.9***
	NMDS1		3.1**	
	Species richness		-2.6**	
	Shannon	2 2*		
Specialist	diversity			
plant species	Jaccard index	3.4***		-3.5***
	NMDS1		-3.7***	
Richness of	Wind	2.6**		-2.9**
plant species	Ant	2.9**		-3.1**
dispersed by:	Dung	3.1**		-3.3***

Table 2. General increase of plant species richness and diversity following inoculation of topsoil removed sites, with the exception of sites at the furthest distances (20 m) from intact remnant grassland patches. Mean (± 1 SE) plant species richness, Shannon diversity and Jaccard index between plots and intact remnant grassland patches, in inoculated and non-inoculated plots, at three distances from the remnant patches. Data for 2012 (prior to inoculation, and 1 year after top soil removal), and for 2013 (one year after the inoculation).

			Inoculur	addition		Distance (m)								
			moculum	audition										
		Inoculated		Non-inc	oculated	:	5	1	0	20				
		2012	2013	2012	2013	2012	2013	2012	2013	2012	2013			
All plant species	Species richness	10.9 ± 1.4	16.2 ± 2.1	11 ± 1.6	12.5 ± 1.3	11.2 ± 1.6	15.3 ± 2	11 ± 1.5	15.2 ± 1.9	10.5 ± 1.4	12.5 ± 1.2			
	Shannon diversity	1.5 ± 0.1	1.7 ± 0.2	1.5 ± 0.1	1.6 ± 0.1	1.5 ± 0.1	1.8 ± 0.1	1.5 ± 0.1	1.7 ± 0.2	1.4 ± 0.1	1.5 ± 0.1			
	Jaccard index	0.4 ± 0.04	0.5 ± 0.04	0.3 ± 0.05	0.4 ± 0.05	0.4 ± 0.04	0.5 ± 0.05	0.4 ± 0.1	0.5 ± 0.05	0.3 ± 0.03	0.3 ± 0.1			
AMF-dependent	Species richness	6.4 ± 0.9	8.4 ± 1.3	6.1 ± 0.8	6.8 ± 0.8	6.5 ± 0.9	8.4 ± 1	6.1 ± 0.9	7.6 ± 0.9	6.2 ± 0.7	6.8 ± 1			
plant species	Shannon diversity	1.2 ± 0.1	1.4 ± 0.1	1.2 ± 0.1	1.2 ± 0.1	1.2 ± 0.1	1.4 ± 0.1	1.2 ± 0.1	1.4 ± 0.1	1.1 ± 0.1	1.1 ± 0.1			
	Jaccard index	0.3 ± 0.03	0.4 ± 0.05	0.3 ± 0.04	0.3 ± 0.03	0.3 ± 0.03	0.4 ± 0.05	0.3 ± 0.03	0.4 ± 0.05	0.3 ± 0.03	0.3 ± 0.02			
Specialist plant	Species richness	2.9 ± 0.4	3.6 ± 0.5	2.8 ± 0.5	3 ± 0.5	3.1 ± 0.4	4 ± 0.5	2.8 ± 0.5	3.9 ± 0.5	2.8 ± 0.5	2.9 ± 0.5			
species	Shannon diversity	0.5 ± 0.1	0.7 ± 0.1	0.5 ± 0.1	0.6 ± 0.1	0.5 ± 0.1	0.8 ± 0.1	0.5 ± 0.1	0.8 ± 0.1	0.4 ± 0.1	0.6 ± 0.1			
	Jaccard index	0.4 ± 0.1	0.6 ± 0.1	0.4 ± 0.1	0.4 ± 0.1	0.4 ± 0.1	0.6 ± 0.1	0.4 ± 0.1	0.6 ± 0.07	0.4 ± 0.05	0.5 ± 0.1			
Richness of	Wind	5.3 ± 0.5	7.2 ± 0.4	5.3 ± 0.6	5.8 ± 0.5	5.3 ± 0.6	6.9 ± 0.5	5.2 ± 0.5	6.7 ± 0.5	5.3 ± 0.5	5.7 ± 0.4			
plant species	Ant	2.2 ± 0.5	3.4 ± 0.4	1.9 ± 0.4	2.2 ± 0.5	2.1 ± 0.4	3.2 ± 0.5	2.1 ± 0.4	3 ± 0.4	1.9 ± 0.4	2.2 ± 0.4			
aloperoed by.	Dung	3.5 ± 0.4	4.8 ± 0.3	3.4 ± 0.4	3.9 ± 0.5	3.3 ± 0.3	4.7 ± 0.4	3.6 ± 0.3	4.6 ± 0.3	3.5 ± 0.4	3.9 ± 0.3			



Figure 1. Spatial arrangement of the study plots relative to the intact grassland

remnants (grey rectangles). Squares represent study plots at 5 m, circles at 10

- m, and triangles at 20 m from the closest intact remnant grassland. Inoculated
- 744 plots are represented by solid symbols and non-inoculated plots by open
- 745 symbols.
- 746



748 Figure 2. Inoculum addition generates strong effects on plant species richness (A), similarity with intact remnant grassland communities (B), and community composition (C) for all plant species, but not in the plots furthest (20 m) from the intact grassland remnants. Inoculated plots: 1; control plots: 0. Error bars represent ± 1 SE. Different lower case letters denote significant differences between years within treatments.



Figure 3. Inoculum addition generates strong positive effects on plant species richness (A), Shannon diversity (B) and similarity with intact remnant grasslands (C) for AMF-dependent plant species, but not in the plots furthest (20 m) from the intact grassland remnants. Inoculated plots: 1; control plots: 0. Error bars present ± 1 SE. Different lower case letters denote significant differences between years within treatments.

- 765
- 766



767 768 Figure 4. Strong effect of inoculum addition on the plant community similarity between plots and intact remnant grassland patches for specialist plant species 769 of nutrient-poor grasslands, but not in the plots furthest (20 m) from the intact 770 771 remnant grasslands. Inoculated plots: 1; control plots: 0. Error bars represent ± 1 SE. Different lower case letters denote significant differences between years 772 773 within treatments.





Figure 5. Strong effects of inoculum addition on the richness of plant species dispersed by ants (A), wind (B) and dung (C), at every distance class, except at 20 m from the intact remnant grassland patches. Inoculated plots: 1, and control plots: 0; Error bars represent \pm 1 SE. Different lower case letters denote significant differences between years within treatments.

- 782
- 783

784 Supplementary material

- 786 **Table S1.** Plant species frequencies in intact remnant grassland patches and in the plots. AMF-dependent plant species,
- specialist plant species of the nutrient-poor grassland and well-dispersed plant species are indicated, and main the seed
- dispersal mode is provided.

	Intact remnant	5 m		10 m		20 m				Dispersal	Well-
Species	grasslands	2012	2013	2012	2013	2012	2013	dependent	Specialist	type	dispersed
Achillea millefolium	2	0	1	0	0	0	0	Х		Wind	
Agrostis canina	2	0	0	1	0	1	0	Х		Wind	
Agrostis capillaris	14	8	14	10	12	0	0	Х	Х	Wind	
Aira caryophyllea	3	0	1	0	0	0	0	Х		Wind	
Anagallis arvensis	7	2	6	2	4	1	2	Х		Wind	
Aphanes australis	7	3	4	2	6	0	3	Х			
Arenaria serpyllifolia	9	7	7	4	7	3	4			Wind	
Artemisia vulgaris	2	0	1	0	0	0	0	Х		Wind	
Betula pendula	11	8	10	7	10	6	8			Wind	
Buddleja davidii	4	1	3	0	0	0	1			Wind	
Calluna vulgaris	15	8	11	7	11	11	13			Wind	х
Campanula rotundifolia	1	1	0	0	0	0	0	Х	Х	Wind	
Cardamine hirsuta	2	1	0	1	0	0	0	Х		Dung	
Carduus crispus	2	0	0	1	0	0	0			Ant	
Carex pilulifera	15	8	11	8	13	12	13		Х	Ant	х
Centaurium erythraea	7	2	6	1	1	1	3	Х		Wind	
Centunculus minimus	2	2	0	0	0	0	0				
Cerastium fontanum	2	2	0	1	0	0	0	X		Dung	
Cerastium glomeratum	1	0	0	1	0	1	0			Dung	

Cirsium arvense	1	0	0	1	0	0	0	Х		Wind	
Conyza canadensis	2	2	0	1	0	1	0	Х		Wind	
Crepis capillaris	2	2	0	0	0	0	0	Х		Wind	
Cirsium spp.	3	0	1	0	3	0	0	Х			
Cytisus scoparius	11	6	7	5	10	2	6	Х	Х	Ant	
Epilobium spp.	1	0	1	0	0	0	0	Х			
Equisetum arvense	3	2	2	2	2	1	1	Х			
Erodium cicutarium	2	1	1	0	1	0	0	Х		Wind	
Fragaria vesca	5	4	4	1	3	0	2	Х		Dung	
Gnaphalium uliginosum	2	2	0	1	0	0	0	Х		Wind	
Holcus lanatus	15	1	12	3	8	0	11	Х		Wind	Х
Holcus mollis	15	3	8	2	8	3	5	Х		Wind	
Hypericum humifusum	3	3	0	3	2	1	2			Wind	
Hypericum perforatum	13	9	12	10	13	7	9	Х		Ant	Х
Hypochaeris radicata	4	1	4	1	4	0	1	Х	Х		
Juncus bufonius	5	0	5	2	4	3	4	Х		Dung	
Juncus effusus	14	12	13	11	14	13	14	Х		Dung	Х
Juncus tenuis	10	6	10	4	7	3	6			Dung	
Linaria vulgaris	3	0	2	0	2	0	3	Х		Wind	
Lotus corniculatus	1	1	1	0	0	0	0	Х	Х	Dung	
Luzula multiflora	15	13	11	14	15	8	13		Х	Dung	Х
Medicago lupulina	6	0	6	0	6	0	1	Х		Dung	
Molinia caerulea	13	5	11	5	9	7	13	Х		Wind	Х
Myosotis spp.	1	0	1	0	0	0	0	Х			
Ornithopus spp.	2	0	2	0	2	0	0		Х		
Pinus sylvestris	15	15	16	14	16	11	16			Wind	X
Plantago major	5	4	3	1	1	1	1	x		Wind	

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Poa annua	9	0	5	2	8	1	6	Х		Wind	
Polygala serpyllifolia	7	0	0	0	0	0	0				
Potentilla erecta	2	0	1	0	0	0	0	Х	Х		
Potentilla reptans	2	1	0	0	0	0	0	Х		Ant	
Prunella vulgaris	5	3	4	0	3	0	1	Х		Ant	
Pteridium aquilinum	4	0	0	1	3	1	2	Х			
Quercus robur	2	0	0	0	1	0	0			Dung	
Ranunculus repens	5	4	3	4	3	0	0	Х		Wind	
Rubus fruticosus agg.	13	11	11	10	12	11	12	Х		Dung	Х
Rumex acetosella	6	0	2	0	5	0	3	Х	Х	Wind	
Sagina procumbens	11	5	9	5	8	2	3			Dung	
Salix spp.	10	8	8	2	5	2	3				
Scrophularia nodosa	4	3	1	1	2	0	1	Х		Wind	
Solidago virgaurea	4	0	3	0	1	0	1	Х		Ant	
Spergula arvensis	1	0	1	0	1	0	0	Х		Dung	
Spergularia rubra	1	0	0	1	0	0	0			Dung	
Stellaria spp.	8	0	7	0	5	0	2	Х			
Taraxacum officinale agg.	5	0	1	0	4	0	0	Х			
Teucrium scorodonia	10	6	5	6	8	5	5	Х		Dung	
Thymus pulegioides	10	0	0	0	0	0	0				
Trifolium dubium	5	1	1	1	1	0	0	Х		Wind	
Vaccinium myrtillus	2	1	0	0	0	0	0			Dung	
Vaccinium spp.	2	0	1	0	1	0	0				
Verbascum thapsus	1	0	0	1	2	0	1	Х		Wind	
Veronica officinalis	8	5	6	5	4	1	3	Х	Х	Ant	
Veronica serpyllifolia	9	3	4	5	7	0	1	X		Ant	
Vicia hirsuta	2	0	1	0	1	0	0	X		Dung	

Asteraceae sp 1	1	0	1	0	0	0	0		
Asteraceae sp 2	1	0	0	0	1	0	0		
Asteraceae sp 3	1	1	0	0	0	0	0		
Brassicaceae sp.	1	0	0	0	0	0	1		
Caryophyllaceae spp.	1	0	1	0	0	0	0		
Lamiaceae spp.	1	0	0	0	1	0	0		
Poaceae spp.	1	0	0	0	0	0	3		