

# Using various artificial soil mixtures to restore dry grasslands in quarries

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1	Using various artificial soil mixtures to restore dry grasslands in quarries
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3	Running head: Substrates for grassland restoration in quarries
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19	

20 Abstract

Quarry restoration is difficult to implement due to the scarcity of the original soil. The restoration 21 22 of Mediterranean dry grasslands to mitigate similar ecosystem/habitat loss is thus still a developing process. So far, very few studies have created artificial substrates to address the lack 23 of soil. In the La Crau plain (southeastern France), quarries were established in an ancient 24 25 Mediterranean steppe-like dry grassland. Once the remaining grassland zones were protected in 26 2001, quarries were extended over former intensive orchards established on the grassland in the 27 1990s before it was protected. Now, restoration has to be done without the unaltered protected 28 grassland soil but with the soil from orchards, which contains fertilizers spread during the orchard exploitation. To recreate a more suitable substrate, the orchard topsoil was mixed with poor 29 30 substrate materials directly extracted from the quarry (0-30 mm pebbles with sandy matrix). Different substrate mixtures were tested with or without sowing the dominant species of the 31 32 grassland, a perennial grass: Brachypodium retusum (and an annual grass Brachypodium hybridum). After five years of monitoring, our results show that raw substrate materials (100%) 33 34 limit the establishment of all species, and that, at the opposite, orchard topsoil (100%) favors a

significantly higher species richness of target and non-target species. The recommended substrate 35 36 mixture to reach a compromise between high target cover and low non-target species cover is composed of 50% raw quarry material and 50% arable soil. Sowing *Brachypodium* reduces target 37 species richness establishment but also limits non-target species richness and cover. 38 39 40 Keywords: Grassland, Raw substrate materials, Mediterranean pseudo-steppe, Artificial soil mixture, Soil restoration, Thero-Brachypodietea. 41 42 43 **Implications for practice:** Soil resources may be scarce for ecological restoration therefore creating substrates based 44 45 on soil mixed with raw quarry material is an opportunity to increase available substrate volumes 46 47 Substrate composed of 100% arable soil maximized both target and non-target species \_ establishment 48 Substrate mixtures composed of 50% arable soil and 50% raw quarry material are a good 49 compromise to create more substrate than there is soil, favor target species and lower 50 non-target species establishment 51 52 -Sowing *Brachypodium* lowers non-target species richness but also target species cover; it creates a physiognomy close to that of the reference grassland. 53

54

#### 55 Introduction

56 Ecological restoration in former quarries is often difficult because of the definitive loss of the original soil resource (Castillejo & Castelló 2010; Jarvis & Walton 2012). Its replacement by low 57 58 quality substrates (e.g. rubble sometimes mixed with inert waste: low nutrient and organic 59 contents, low mycorrhize, poor soil structure, low water storage, etc.) is often insufficient for 60 successful revegetation because it limits the establishment and growth of plants, which are also 61 exposed to recurrent drought stress in the Mediterranean region (Oliveira et al. 2011). To date, very few studies have focused on the creation in integrum of a soil in ecological restoration 62 projects, while this is known to be one of the key drivers of restoration success (Mchergui et al. 63 2014; Muñoz-Rojas 2018). 64

When quarrying is performed, soils can be discarded or stockpiled, altering their quality (Stahl et al. 2002). If quarrying occurs in pristine / undamaged ecosystems, immediately transferring the soil to a restoration site (just-in-time soil transfer) is a good way to restore high species-diversity grasslands in Southeastern Mediterranean France (Jaunatre et al. 2014; Bulot et

al. 2017; Buisson et al. 2018). However, this technique is hardly applied because it is costly and 69 70 need to be performed concomitantly with undesirable soil extraction on undamaged well preserved ecosystems. It is thus limited to cases where a reference site is planned to be destroyed 71 72 (Bullock 1998).. Other sources of soils, such as from abandoned agricultural areas may also be available for restoration. However, fertilizers spread in the course of intensive exploitation 73 (nitrogen, phosphorus and potassium) may have considerably increased the nutrient contents of 74 75 these soils (Römermann et al. 2005; Wang et al. 2017; Helm et al. 2019). The use of the arable 76 layer (i.e. topsoil) resulting from abandoned agricultural areas might therefore favor the 77 establishment of undesirable species from the seedbank (Buisson et al. 2018), including ruderal 78 and exotic invasive species, to the detriment of target stress-tolerant plant species (Castillejo & 79 Castelló 2010; Ballesteros et al. 2012; Jaunatre et al. 2014). Thus, the use of such arable soil may not enable the restoration of nutrient poor grassland vegetation with high species-diversity, thus 80 81 impeding the restoration of such ecosystems.

The selection of the materials at the outset of a project determines the success of the restoration process (Bradshaw 2000), and is particularly decisive for the recovery of the target vegetation associated with specific substrates (Ballesteros et al. 2014, 2012; Whiting et al. 2004). In order to recreate an adequate substrate starting with arable soil as a base, mixing it with natural materials (for example, originating from the geological substrate), not impacted by the chemical inputs of former agricultural practices, should allow approaching the oligotrophic characteristics of the soils of the reference communities.

The first aim of the present experimental setup was to test different mixtures of substrates 89 in the context of a Mediterranean dry grassland (La Crau, Southern France) restoration. This will 90 allow us to select the best substrate for initiating the successional trajectory of the vegetation 91 towards the reference dry grassland (Asphodeletum fistulosi - Molinier & Tallon 1950). During 92 93 the restoration process, species may not recolonize owing to a lack of seed dispersal. In that case, 94 the recolonization process may be accelerated by sowing target species, or by sowing so-called 'nurse' species to encourage the establishment of other species (Padilla & Pugnaire 2006; Ren et 95 96 al. 2008; López & Cavallero 2017). The second aim of this experimental setup is thus to test whether Brachypodium retusum (Poaceae) can act as a nurse species for grassland restoration in 97 98 quarries. It is a perennial rhizomatous grass species which represents the most abundant species 99 in biomass and cover in many north-western Mediterranean grasslands (Molinier & Tallon 1950; 100 Luis et al. 2004; Buisson et al. 2021). It is common within its natural range (Contu 2013), where it plays an important role particularly in post-wildfire recovery (Caturla et al. 2000). B. retusum 101 102 is one of the first species to reproduce after a fire (Vidaller et al. 2019), and to rapidly recolonize

the environment. It plays a key role in protecting the soil against erosion, favoring the establishment of target shrub species and preventing the establishment of undesirable species (Caturla et al. 2000; Larchevêque et al. 2010; Raventós et al. 2012). Its rather tall and dense physiognomy can be expected to enable the capture of windborne seeds (Bullock & Moy 2004; Giladi et al. 2013) and the protection of seedlings from grazing. Consequently, *B. retusum* would be a good candidate to have an indirect positive impact on the establishment and survival of target species (defined here as the species found in the reference grassland).

The experimental setup was designed to test the effect of (1) the physical-chemical 110 111 composition of the substrate, (2) B. retusum sowing and (3) their interaction, on the natural colonization of the regional species pool. We tested the following hypotheses: (1) The substrate 112 113 mixtures, constituted exclusively of materials extracted from the quarry (pebbles of 0-30 mm with sandy matrix hereafter raw quarry material), are poor in nutrients and organic matter, and 114 115 will limit the establishment of all species, i.e. undesirable ruderal species and target species; (2) Spontaneous colonization enables plant communities to tend towards the reference grassland, as 116 117 long as i) they are not in competition with dense patches of ruderal species (like on the substrate mixtures with the highest proportion of arable soil), or ii) the abiotic conditions are not too harsh 118 119 (like on the substrate mixtures with raw quarry material); and (3) Sowing Brachypodium retusum enables the rapid establishment of this species, and potentially facilitates the trapping of 120 windborne seeds from the reference grassland. Sown plots will then have higher cover of B. 121 *retusum*, increased richness of target species and greater similarity to the reference grassland. 122

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#### 124 Methods

125 *Study area* 

Our study was carried out on a quarry in the La Crau plain, located in southeastern Mediterranean 126 France (Bouches-du-Rhône). An area of the gravel quarry "La Ménudelle" was dedicated to this 127 study. It was in operation from 2008 to 2011, impermeabilized in 2012 (using deposits of 128 decantation sludge - clays and silt - presenting a strong water retention capability) and filled up 129 130 in 2012-2013 (with inert materials that are non-fermentable, non-polluting and not likely to alter the physical and chemical properties of the soil down to 3.50m depth (BO-MEDDTL 2011; 131 132 Figure 1). After compacting and levelling of the surface of the landfill, the site was then at an elevation of 40 cm below the original surface of the reference grassland and ready to receive the 133 134 experimental setup.

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136 *The different substrate mixtures* 

Two types of substrates were used in the experiment for the different mixtures. 1) Arable soil 137 from an orchard abandoned in 2015, composed of the topsoil (organo-mineral layer, 0-20 cm) 138 and the subsoil (mineral layer, 20-40 cm) of cultivated Haplic Cambisol soil mixed together 139 (Chenot et al. 2018). It was directly collected at the donor site at the same time as the different 140 treatments were constituted to avoid any storage period of the soil and any enrichment in 141 undesirable non-target species during storage (in particular windborne Asteraceae species) 142 (Bordez 2015). 2) The raw quarry material was composed of sand and of alluvial gravel, sifted 143 so as to keep only elements between 0 and 30 mm, also stored for less than 24h as in order to 144 145 avoid enrichment with seeds of non-target species. Overall, five substrate mixtures with different 146 proportions of each substrate were tested : 100% raw quarry material, 75% raw quarry material 147 with 25% arable soil, 50% raw quarry material with 50% arable soil, 25% raw quarry material with 75% arable soil, and 100% arable soil (Table 1). 148

The various mixtures made it possible to obtain significant differences in substrate mixture physical-chemical properties (Supplement S1). The mixtures were spread in spring 2016 in the form of 30-40 cm thick 5m × 5m plots (similar thickness of soil to the reference ecosystem) using back hoes and hand tools. During the mixing of materials using backhoes, the soil was stirred, mixed and thus aired to ensure the growth of the plants (Jarvis & Walton 2012). Overall, 50 plots were installed at the site in February 2016 (Supplement S2).

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#### 156 *Sowing the dominant species*

Two sowing treatments were tested: (1) Each substrate mixture without sowing of any species 157 158 (natural colonization); (2) Each substrate mixture with sowing of Brachypodium spp.. Commercially available non-certified seeds of non-local origin (Spain) were sown (germination 159 rate: 73%) in March 2016 over  $625m^2$  of experimental plots ( $25m^2 \times 25$  plots), with a ratio of 160 10g of seeds/m<sup>2</sup> (3000 seeds/m<sup>2</sup>). While we purchased *Brachypodium retusum* seeds only, our 161 seed mix turned out to be only 32% B. retusum and 68% Brachypodium hybridum, the latter 162 being an annual Brachypodium which seems to have been confused during seed collection by the 163 164 seed company. However this species is also present in the reference grassland with a lower plant cover (Pavon & Pires 2020). Throughout the remainder of this article, only the term 165 Brachypodium was used for the sown seed mix. The experimental setup thus includes five 166 replicates of each treatment (5 substrate mixtures  $\times$  2 sowing treatments  $\times$  5 replicates). 167

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169 Experimental implementation

The different treatments were spread randomly over the site (Supplement S2). No material was 170 171 spread in the paths between the plots. Once a year, paths were mowed using a grass trimmer to limit their colonization by plants and avoid their seeding, which might have influenced the 172 establishment of species on the plots by edge effects. As the reference grassland is grazed 173 (Saatkamp et al. 2010; Saatkamp et al. 2018), grazing by an itinerant flock of sheep, connecting 174 the reference grassland and all the plots, was reintroduced after the experiment was set up (for a 175 total of 3 days a year between February and May). Finally, a drainage outlet was dug at the end 176 of the experimental plot to avoid the stagnation of water during rainy episodes. 177

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#### 179 Annual vegetation surveys

180 Monitoring of the vegetation was carried out every year from 2016 to 2018 and in 2020 to assess plant recolonization dynamics and to compare the different treatments. For that purpose, 181 182 permanent 4m<sup>2</sup> quadrats situated at the center of each 25m<sup>2</sup> plot (to avoid any edge effect) were monitored in May 2016, May 2017, May 2018 and May 2020. The abundance of each plant 183 184 species was estimated on the basis of percent cover (Gillet 2000) and total plant cover was also estimated. The vegetation developing on the quadrats was compared with the reference 185 ecosystem after a survey of randomly placed 4m<sup>2</sup> quadrats on the reference grassland: ten 186 187 quadrats in 2018 and five quadrats in 2020. Reference plant communities contained 52.2±2.2 species per 4m<sup>2</sup>, and 109 species overall in these 15 quadrats. 188

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190 Data analyses

All the data analyses were performed using the packages *vegan* (Oksanen et al. 2015), *car* (Fox
et al. 2020), *multcomp* (Hothorn et al. 2020), *emmeans* (Lenth et al. 2020) and *lme4* (Bates et al.
2016) in R 4.0.3. (R Core Team 2020).

194 We evaluated the effects of substrate mixtures, sowing and time on species composition, richness and cover of plant communities. We also evaluated the effects of substrate mixtures and 195 sowing five years after the experiment was set up, in 2020. This was done on species 196 197 composition, richness and cover of plant communities, as well as on *Brachypodium retusum* and Brachypodium hybridum covers. In 2020, we also assessed the effects of treatments on plant 198 199 species richness and plant cover for target and non-target species separately (target species being the species found on the reference grassland plots in 2020). For species composition, we used the 200 201 data of all species but Brachypodium spp. as the aim was to assess the effect on the plant community independently of *Brachypodium* cover. As the sum of species cover usually largely 202 203 exceed total species cover, we therefore used as cover without *Brachypodium* the product of the

sum of species cover excluding *Brachypodium* spp. cover by the estimated total plant cover,
divided by the sum of all species cover, hereafter called plant cover without *Brachypodium*.

Two indices were used to assess the state of the plant communities compared to the 206 207 reference ecosystem. 1) The CSII<sub>norm</sub> Index (normalized Community Structure Integrity Index) 208 measuring the proportion of the abundance of species of the reference community represented in the restored community (it shows the percentage of similarity in terms of composition and 209 abundance of the communities studied compared to the reference community). 2) The HAI Index 210 (Higher Abundance Index) measuring the proportion of the abundance of species in the restored 211 212 community which is higher than in the reference community (it shows what is 'in excess', in terms of species and abundance in the studied communities studied compared to the reference 213 214 community) (Jaunatre et al. 2013).

In order to characterize the composition of the plant communities sampled on the different plots, two Non-Metric Multidimensional Scaling (NMDS) based on Bray–Curtis similarity were performed (Borcard et al. 2011). One NMDS was run on all years (200 quadrats (5 mixtures  $\times$  2 sowing treatments  $\times$  5 replicates  $\times$  4 years (2016, 2017, 2018, 2020))  $\times$  155 species) and one in 2020 (55 quadrats (5 mixtures  $\times$  2 treatments  $\times$  5 replicates + 5 quadrats in the reference grassland)  $\times$  135 species).

To test the effects of the substrate mixtures  $\times$  *Brachypodium* sowing  $\times$  time on plant species richness, plant cover, CSII<sub>norm</sub> and HAI, one Generalized Linear Mixed Model (GLMM) was performed on each of these variables. For each, we used substrate mixtures, *Brachypodium* sowing and years as fixed factors and quadrats as a random factor to take repeated measures into account. They were followed by pairwise contrast comparisons with a Tukey adjustment when significant.

To test for possible effects of the substrate mixtures  $\times$  *Brachypodium* sowing on plant species richness, plant cover CSII<sub>norm</sub> and HAI in 2020, one Generalized Linear Model GLM was performed on each of these variables. They were followed by pairwise contrast comparisons with a Tukey adjustment when significant. All models were fitted with a poisson error distribution for plant species richness and a gamma error distribution with log link for the other variables.

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#### 234 **Results**

235 Effects of substrate mixtures, Brachypodium sowing and time on plant communities

All plant communities changed over time (Supplement S3). They all started in 2016 with arable
weeds, such as *Diplotaxis tenuifolia* (L.) DC., *Polygonum aviculare* L., *Solanum nigrum* L. and *Veronica persica* Poir.. With time, they gained target species of the reference grassland, such as *Avena barbata* Pott ex Link, *Catapodium rigidum* (L.) C.E.Hubb., *Hordeum murinum* L., *Medicago truncatula* Gaertn., *Rostraria cristata* (L.) Tzvelev, *Trifolium campestre* Schreb., *Trifolium glomeratum* L. and *Trifolium scabrum* L..

Overall, species richness increased with time on all plots. It increased by 5 fold between 2016 and 2017 and only by 10% from 2017 to 2020 (Supplement S3). Percent covers of unsown species increased by two fold from 2016 to 2018, then decreased by 10% in 2020 (Supplement S3). Substrate mixtures with 100% or 75% of raw quarry material had significantly lower species richness and lower percent cover of unsown species than other substrate mixtures on all years. Plots where *Brachypodium* was not sown had significantly higher species richness and higher percent cover of unsown species (Supplement S3).

CSIInorm increased by ten fold between 2016 and 2017 (especially on unsown plots and on sown plots with the highest proportion of arable soil) and then remained relatively stable or underwent a 20% decrease by 2020 (Supplement S3). HAI on unsown plots were always high (close to 1). HAI on sown plots first increased by 3 fold and then decrease by the same order to go back below 0.6 in 2020 (Supplement S3).

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#### 255 Effects of substrate mixtures on Brachypodium retusum cover in 2020

*Brachypodium retusum* had a significantly higher cover on sown plots ( $\chi^2=381.23$ , df=1 p<0.001\*\*\*; Figure 2). It established very well on all sown plots (found on 25 plots out of 25 sown in 2020; average percent cover = 38.8%). It was found on two of the 25 unsown plots in 2020, but with a significantly lower percent cover on average (0.04%). *Brachypodium hybridum* had a significantly higher cover only on plots sown on the substrate mixture with 100% arable soil ( $\chi^2=23.97$ , df=4, p<0.001; Figure 2). Its percent cover (8.8%) was much lower than that of *Brachypodium retusum* (35%).

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#### 264 Effects of substrate mixtures and Brachypodium sowing on plant communities in 2020

*Brachypodium* sowing and the substrate mixtures influenced plant community composition in 2020 (Figure 3). Axis 1 of the NMDS on plant community data showed two major groups: the quadrats where *Brachypodium* was sown on the left and those where *Brachypodium* was not sown on the right. The sown quadrats were characterized by target species such as *Andryala* 

integrifolia L., Lobularia maritima (L.) Desv. and Medicago monspeliaca (L.) Trautv.. In

contrast, those without sowing of *Brachypodium* were characterized by *Dittrichia viscosa* (L.)
Greuter, *Oloptum miliaceum* (L.) Röser & Hamasha and *Symphyotrichum squamatum* (Spreng.)
G.L.Nesom. Axis 2 of the NMDS showed a slight gradient in the structure of the plant
communities in relation to substrate mixtures, and the reference grassland (Figure 3): mixtures
with higher proportions of arable soil sown with *Brachypodium* were slightly more similar to the
reference grassland than mixtures with higher proportions of raw quarry material.

Substrate mixtures with a higher proportion of arable soil had a significantly higher 276 species richness of both target and non-target species (respectively:  $\chi^2=37.94$ , df=4, p<0.001 and 277  $\chi^2$ =30.94, df=4, p<0.001, Figure 4). *Brachypodium* sowing did not influence target plant species 278 richness ( $\chi^2=0.05$ , df=1, p=0.825), and plots where *Brachypodium* was sown had lower richness 279 of non-target species ( $\chi^2$ =13.10, df=1, p<0.001, Figure 4). Substrate mixtures with a higher 280 281 proportion of arable soil also had a higher plant cover of target species, although this was 282 significant only on plots sown with *Brachypodium* ( $\chi^2$ =38.57, df=4, p<0.001, Figure 5). The plant cover of non-target species was only significantly higher on unsown plots with a substrate 283 284 mixture with 100% of arable soil compared with sown plots with the same substrate mixture  $(\gamma^2 = 13.62, df = 4, p = 0.009, Figure 5).$ 285

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287 Similarity of plant communities obtained in the different soil mixtures to those of the reference288 grassland

In 2020, 60 species, out of the 85 species inventoried in the five reference grassland plots, were 289 recorded on the experimental substrate mixtures (50 plots - total number of species in 2020 = 290 136). Among these 60 species, we found Anisantha madritensis (L.) Nevski, Anisantha rubens 291 (L.) Nevski, Catapodium rigidum (L.) C.E.Hubb., Crepis foetida L., Galactites tomentosus 292 Moench, Medicago monspeliaca (L.) Trautv., Rostraria cristata (L.) Tzvelev, Sonchus oleraceus 293 L., Trifolium campestre Schreb., Trifolium scabrum L. and Vulpia ciliata Dumort. (Supplement 294 295 S4, S5). The plant communities developing on newly created substrates remained highly different from the reference grassland five years after implementation (the best CSIInorm reached 18.15% 296 for substrates composed of 100% arable soil;  $\chi^2=31.08$ , df=4, p<0.001; Figure 6). The substrate 297 mixtures with the poorest nutrient content (QM and QM\_AS), presented the plant communities 298 299 that were farthest from the reference grassland in terms of composition and abundance, in contrast to those with a high proportion of arable soil, such as AS and AS\_QM (Figure 6). The 300 CSIInorm restoration index also showed that plots where Brachypodium was not sown were more 301 similar to the grassland than where *Brachypodium* was sown ( $\chi^2$ =11.89, df=1, p<0.001; Figure 302 303 6).

Unsown plots contained species not found in the reference grassland nor on sown plots, 304 305 such as Artemisia verlotiorum Lamotte, Clinopodium nepeta (L.) Kuntze and Bituminaria bituminosa (L.) C.H.Stirt.. Sown plots contained species not found in the reference grassland nor 306 307 on unsown plots, such as *Hirschfeldia incana* (L.) Lagr.-Foss. and *Medicago orbicularis* (L.) Bartal. (Supplement S4, S5). Unsown plots contained 10 reference grassland species not found 308 309 on sown plots, such as Carduus nigrescens Vill., Eryngium campestre L., Linum strictum L., Neatostema apulum (L.) I.M.Johnst., Sideritis romana L. and Urospermum dalechampii (L.) 310 Scop. ex F.W.Schmidt. Sown plots contained 6 reference grassland species not found on unsown 311 312 plots, such as Aira cupaniana Guss., Logfia gallica (L.) Coss. & Germ., Petrorhagia prolifera 313 (L.) P.W.Ball & Heywood and Plantago berlardii All. (Supplement S4, S5). Most plots also 314 contained high abundances (>5%) of species that are common in the reference grasslands but usually recorded there with lower abundances (<0.5%) (A. madritensis, R. cristata, H. murinum, 315 316 A. rubens, T. campestre, A. barbata, M. truncatula, C. foetida, Geranium molle, etc.). These three sets of species explain the relatively high HAI index (86.4%; Figure 6; Supplement S4, S5). The 317 318 higher HAI restoration index on the unsown plots showed that sowing Brachypodium limited the establishment of non-target species, but this was significant only on substrate with 100% raw 319 320 quarry material ( $\chi^2$ =11.89, df=1, p<0.001; Figure 6). There was no significant differences of HAI between substrate mixtures ( $\chi^2$ =2.42, df=4, p=0.660; Figure 6). 321

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#### 324 Discussion

Despite the fact that plant communities are still distant from the reference grassland after five 325 years, the tested substrate mixtures allow the rapid establishment of a grassland-like vegetation 326 with numerous Poaceae species. The total number of target species (defined here as the species 327 328 found in the reference grassland) increased with the age of succession in the majority of substrate mixtures as already demonstrated in other restoration experiments undertaken in the La Crau 329 grasslands (Coiffait-Gombault et al. 2012; Bulot et al. 2014) or in other dry grasslands (Zobel et 330 331 al. 1996). The first years after restoration are often characterized by weedy ruderal species (Vida et al. 2010). Our results confirmed this pattern with a strong dominance of ruderal species, such 332 333 as Diplotaxis tenuifolia, Polygonum aviculare, etc. which then become rare or are replaced by a mix of ruderal-competitive species sensu Grime (1977) and some target species all of which 334 naturally colonize the site from the available regional pool of species. The most common 335 reference grassland species found on the experimental plots after five years are i) wind-dispersed 336 337 Asteraceae (Crepis foetida, Carduus pycnocephalus, Galactites tomentosa), ii) small Poaceae

(Anisantha madritensis, Anisantha rubens, Catapodium rigidum) which are common in the short-338 term seed bank and can easily be transported by wind over small distances as well as by sheep 339 and ants, and iii) Fabaceae (Trifolium campestre, Trifolium glomeratum, Trifolium scabrum, 340 Medicago truncatula) which are found in the short-term and long-term seed banks and are 341 transported by ants (Buisson et al. 2006; De Almeida et al. 2020). This finding is similar to that 342 of Alday et al. (2011), where species that exhibited zoochory and anemochory were most 343 abundant in the early stages of post-mining succession. Monitoring over the longest term possible 344 345 is recommended as recovery rates and patterns of change in the plant communities may be slow 346 (Jones et al. 2018). Recovery is particularly slow in the La Crau grasslands (Coiffait-Gombault et al. 2012; Bulot et al. 2014). In the long term, it is expected that the plant communities on the 347 348 different substrate mixtures will keep developing in different directions and will never achieve the entire restoration of the reference grassland because of highly different starting conditions 349 350 (substrate mixtures) (Prach et al. 2014). Such results have been already demonstrated for La Crau 351 grasslands with former cultivation (Jaunatre et al. 2016) and sheep paddocks (Saatkamp et al. 352 2020) which still show different plant composition and diversity to the reference grassland several centuries or millennia after their abandonment (Römermann et al. 2005; Helm et al. 353 354 2018).

While some target species colonized the restored areas, the high Higher Abundance 355 Indices (HAI) show that some target species over develop (compared to their abundance on the 356 reference grassland) and that non-target species are common. The latter may be due to the fact 357 that the site is not directly surrounded by reference grassland areas, which impede possibilities 358 359 of target species recolonization (Lanta et al. 2020). The neighboring areas are strongly impacted by quarry exploitation, former quarries, or former orchards where ruderal species predominate. 360 Non-target species may thus easily find their way to the experimental plots, carried in the seed 361 362 bank of the arable soil used for the experiment, by the wind or by itinerant grazing, sheep transporting indistinctively target and non-target species (Albert et al. 2015). The pool of plant 363 species, that is, the source of seeds, of diaspores and the potential for colonization of species 364 plays a major role in the restoration, by influencing the process of spontaneous succession 365 (Řehounková & Prach 2008; Strykstra et al. 1998; Zobel et al. 1998). Therefore, the 366 367 characteristics of the landscape should be properly taken into account for any restoration initiatives (Bakker et al. 1998; Prach et al. 2001; Son et al. 2020), and in our case, may have a 368 369 higher success if adjacent to patches of the reference ecosystem (Buisson et al. 2006).

The substrates represent major variables determining the rhythm and direction of plant 371 372 successions and should unavoidably be taken into account in restoration programs (Prach et al. 2001). The substrate mixtures poor in nutrients, i.e. with a higher percentage of raw quarry 373 374 material, may not be the best as they have lower target species plant covers and lower species 375 richness of both target and non-target plant species than the richer soils (with a higher percentage of arable soil). Survival of target species is indeed severely impeded by harsh abiotic conditions 376 (Pueyo et al. 2009). Moreover, as they generally have a lower plant cover, these substrate 377 378 mixtures may lead to higher soil erosion, the region having very strong winds blowing 334 379 days/year (Buisson & Dutoit 2004).

380 In the present study, fertilizers (N, P, K) applied during the phase of exploitation of the 381 orchards (1983-2015) and thus found in the arable soil have certainly contributed to increasing plant cover on richer substrate mixtures, as it has already been demonstrated elsewhere (Norman 382 383 et al. 2006). Higher plant species richness on nutrient-richer soils is likely due to the fact that arable soils contain a seed bank while raw material does not (Bulot et al. 2014). Substrate 384 385 mixtures with 50% or less of arable soils have significantly less non-target species richness and substrate mixtures with 50% or more of arable soils have significantly more target species 386 387 richness or cover. Therefore, in order to favor target species richness and cover and maintain non-target cover as low as possible, the best compromise would be the substrate mixture with 388 50% arable soil and 50% raw quarry material. This substrate mixture also has a similarity to the 389 reference grassland plant community which is amongst the highest. 390

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Sowing Brachypodium seeds allowed Brachypodium retusum to establish on all plots, 392 although its cover was lower on the raw quarry material. In 2020, sown plots have a lower 393 richness of non-target species, a lower cover of target species, and a lower similarity to the 394 395 reference grassland (as well as a lower cover of non-target species on arable soils only). These 396 results contrast with other studies where seeding increased the establishment of target species (Baasch et al. 2012; Norman et al. 2006). These experiments were however greatly different from 397 398 this one: we sowed one dominant grassland species and they sowed species-rich mixtures or hay collected from a reference grassland. B. retusum therefore seems to compete with non-target 399 400 species establishment and with target species growth: its clonal growth and ability to cover the ground, and its rhizome and root system make it a quite competitive species (Bonet 2004; Clary 401 et al. 2004; Saatkamp et al. 2018). This seems to contradict Raventós et al. (2012) study which 402 showed a facilitative effect of *B. retusum* during the establishment phase of target species after 403 404 fire. However, that study was carried out on shrubby chamaephyte species (Ulex parviflorus

405 Pourr., *Cistus albidus* L., *Helianthemum marifolium* Mill. and *Ononis fruticosa* L.) and not on 406 grassland plants as it is the case here. The reason for a greater competitive effect of *B. retusum* 407 on target species than on non-target species cover most probably lays in the fact that target 408 grassland species are stress tolerant and non-target species rather ruderal (e.g. *D. tenuifolia*) and 409 competitive (e.g. *O. miliaceum*) species.

There was no significant effect of *B. retusum* on target species richness. We can therefore 410 conclude that B. retusum impacts target species cover rather than target species establishment. In 411 addition, these data suggest that B. retusum may not facilitate target species arrival by capturing 412 413 windborne seeds as we initially hypothesized, although this could also be due to the fact that few 414 target species seeds were wind-blown onto the experimental plots as there is no adjacent 415 reference grassland nearby. While we show that B. retusum cover is much higher than B. hybridum, it is not possible to fully disentangle the specific effect of B. retusum from that of B. 416 417 hybridum. However, so far, it is still not possible to obtain pure and certified seeds of B. retusum in Southern France. Future restoration projects in Southern France will therefore still use this 418 419 mixture of *B. retusum* + *B. hybridum* from Spain, until a better option is available for purchase. 420

421 While plots with a substrate composed of 100% of arable soil and not sown with 422 Brachypodium had the highest richness of target species and highest similarity of the plant communities to the reference grassland community, they also had the highest cover of non-target 423 species. Moreover, using a substrate composed of 100% arable soil may limit opportunities for 424 restoration activities due to insufficient substrate being available to restore more surface area 425 426 than the area from which it was excavated. In order to increase the volume of substrate produced 427 to restore wider areas than solely that of abandoned orchards (but also quarries), arable soil should be mixed / diluted with raw quarry material. Therefore, the next best option would be the 428 substrate mixture with 50% arable soil and 50% quarry material. This substrate mixture does not 429 430 differ significantly from the 100% arable soil mixtures in similarity to the reference grassland community and target species richness and cover. It is then a good compromise for increasing 431 432 target and decreasing non-target species richness and cover.

Not sowing *Brachypodium* allows better growth and thus higher cover of target species; but also leads to higher non-target species richness and cover and some over-abundance of target species (*A. madritensis, R. cristata, H. murinum, A. rubens, T. campestre, A. barbata, M. truncatula, C. foetida, Geranium molle*, etc.). However, sowing *Brachypodium*, while not preventing the colonization of non-target species, does reduce their cover and richness and creates a physiognomy similar to that of the reference grassland with *B. retusum* as a dominant grass. *B. retusum* is known to be extremely slow at recolonizing areas with disturbed soils
(Coiffait-Gombault et al. 2012). Hence, sowing *B. retusum* is a major step towards reintroducing
it.

Further studies should look at enriching sown and unsown plots with a few target species to identify the potential key roles of *B. retusum* in interacting with grassland species establishment. A new experiment could be about sowing a more diverse seed mixture of target species (which is still a problem as most seeds of the target species are not available for purchase) and whether sowing of *B. retusum* and this seed mix should be simultaneous or sequential. Longer monitoring of these experimental plots may allow us to refine the best treatment to be applied, i.e. where non-target species decrease the most and where target species increase.

449 450

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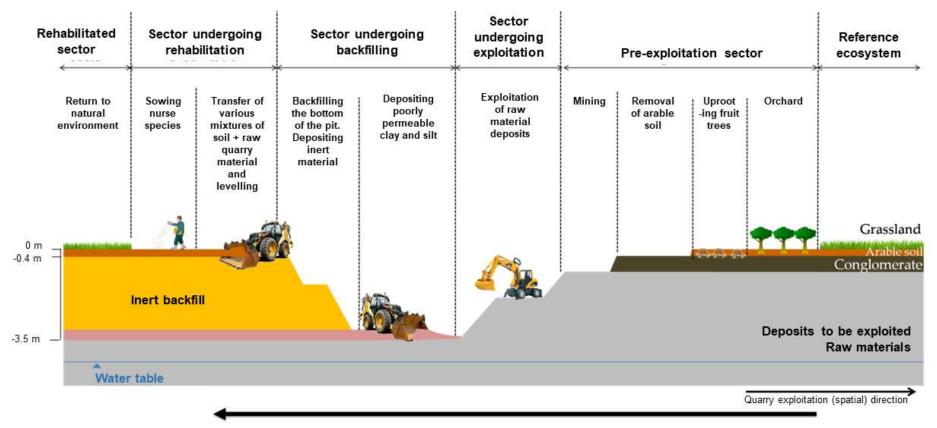
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**Table 1**: Composition of the different mixtures used for the experiment.

## 

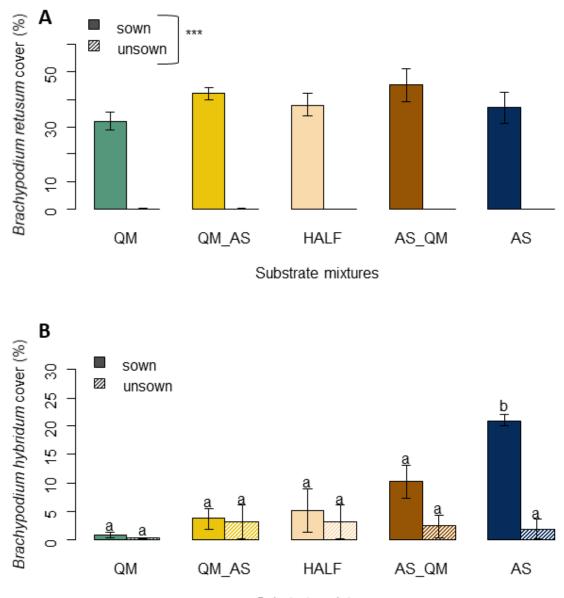
Code for	raw Quarry Material	Arable Soil	Color code
figures	(QM)	(AS)	for figures
QM	100%	0%	
QM_AS	75%	25%	
HALF	50%	50%	
AS_QM	25%	75%	
AS	0%	100%	



Quarry exploitation and restoration (temporal) direction

661662 Figure 1. Diagram of the exploitation and restoration of the Ménudelle quarry in Southeastern France.

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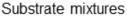
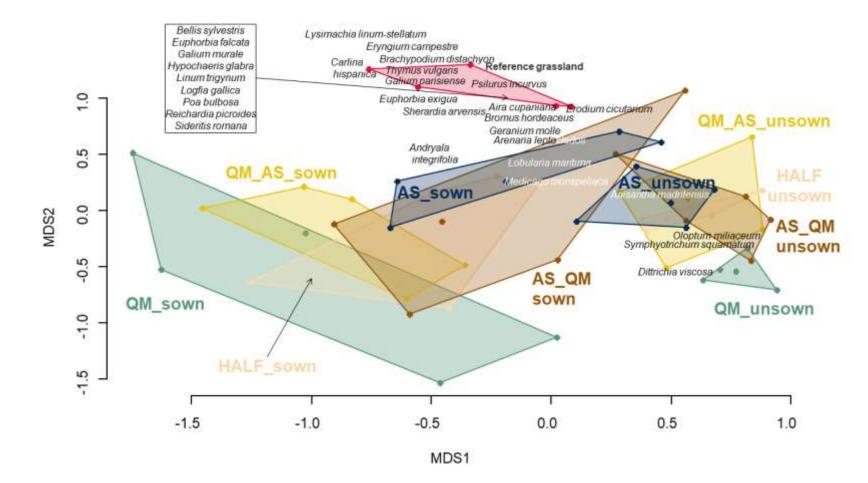


Figure 2. Brachypodium cover on experimental plots in 2020 (mean cover ± standard error). Plots 667 668 were sown (dark grey) or not (light grey) with Brachypodium. Substrate mixtures included QM - Quarry raw Material or AS - Arable Soil (see Table 1 for mixture codes). A. Brachypodium 669 670 retusum (mean cover = 38.8%): Substrate mixtures  $\chi^2$ =4.98, df=4, p=0.290 n.s.; Brachypodium sowing  $\chi^2$ =381.23, df=1, p<0.001\*\*\* (Substrate mixtures × *Brachypodium* sowing  $\chi^2$ =5.05, df=4, 671 p=0.282 n.s.). B. Brachypodium hybridum (mean cover = 8.2%): Substrate mixtures  $\times$ 672 Brachypodium sowing  $\chi^2=23.97$ , df=4, p<0.001\*\*\* (Substrate mixtures  $\chi^2=25.07$ , df=4, 673  $p<0.001^{***}$ ; Brachypodium sowing  $\chi^2=17.44$ , df=1,  $p<0.001^{***}$ ). 674



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Figure 3. Plant species composition in 2020 using NMDS. Polygons indicate of plots with the same treatment (NMDS stress: 0.21). Plots were
sown (sown) or not (unsown) with *Brachypodium*. Substrate mixtures included QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry
Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil
(see Table 1 for mixture codes).

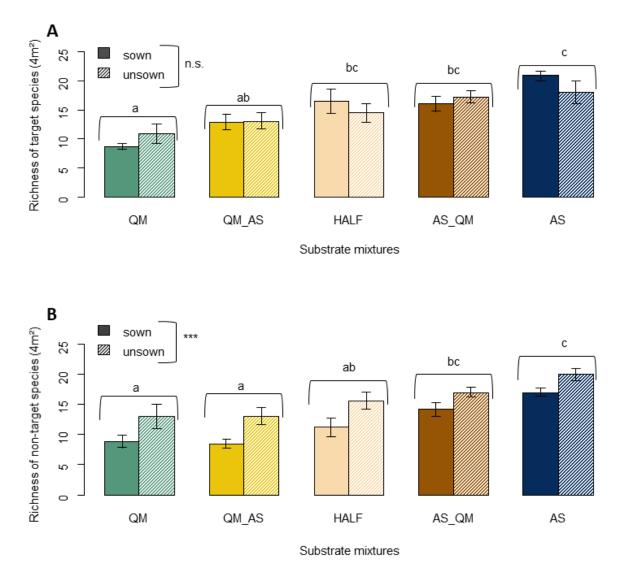


Figure 4. Mean species richness (± standard error) of A. target species and B. non-target species on experimental plots in 2020. Plots were sown (dark grey) or not (light grey) with *Brachypodium*. Substrate mixtures included QM – Quarry raw Material or AS – Arable Soil (see Table 1 for mixture codes). A. Substrate mixtures  $\chi^2=37.94$ , df=4, p<0.001\*\*\*; *Brachypodium* sowing  $\chi^2=0.05$ , df=1, p=0.825 n.s. (Substrate mixtures × *Brachypodium* sowing  $\chi^2=3.09$ , df=4, p=0.543 n.s.). B. Substrate mixtures  $\chi^2=30.94$ , df=4, p<0.001\*\*\*; *Brachypodium* sowing  $\chi^2=13.10$ , df=1, p<0.001\*\*\* (Substrate mixtures × *Brachypodium* sowing  $\chi^2=2.06$ , df=4, p=0.726 n.s.).

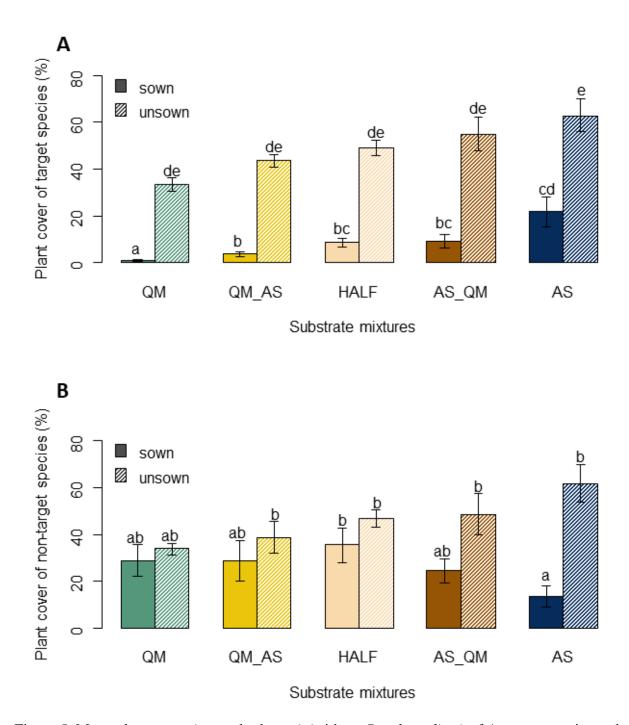


Figure 5. Mean plant cover (± standard error) (without *Brachypodium*) of A. target species and B. non-target species on experimental plots in 2020. Plots were sown (dark grey) or not (light grey) with *Brachypodium*. Substrate mixtures included QM – Quarry raw Material or AS – Arable Soil (see Table 1 for mixture codes). A. Substrate mixtures × *Brachypodium* sowing  $\chi^2$ =38.57, df=4, p<0.001\*\*\* (Substrate mixtures  $\chi^2$ =61.00, df=4, p<0.001\*\*\*; *Brachypodium* sowing  $\chi^2$ =174.19, df=1, p<0.001\*\*\*). B. Substrate mixtures × *Brachypodium* sowing  $\chi^2$ =13.62, df=4, p=0.009\*\* (Substrate mixtures  $\chi^2$ =2.01, df=4, p=0.735 n.s.; *Brachypodium* sowing  $\chi^2$ =17.81, df=1, p<0.001\*\*\*).

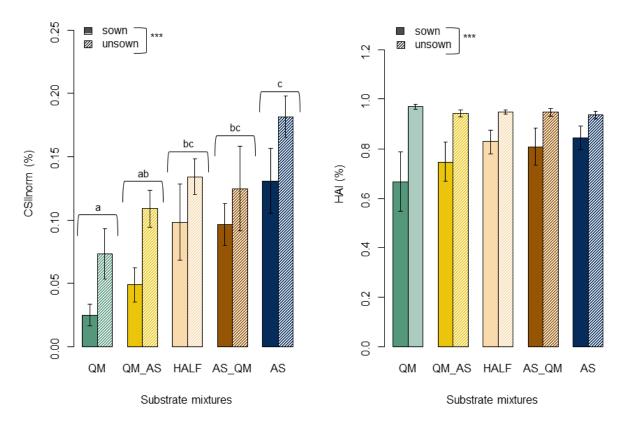


Figure 6. Mean CSII and HAI (± standard error) on experimental plots in 2020. A. CSII: Substrate mixtures  $\chi^2=31.08$ , df=4, p<0.001\*\*\*; *Brachypodium* sowing  $\chi^2=11.89$ , df=1, p<0.001\*\*\* (Substrate mixtures × *Brachypodium* sowing  $\chi^2=0.815$ , df=4, p=0. 936 n.s.). B. HAI: Substrate mixtures  $\chi^2=2.42$ , df=4, p=0.660; *Brachypodium* sowing  $\chi^2=19.98$ , df=1, p<0.001\*\*\* (Substrate mixtures × *Brachypodium* sowing  $\chi^2=4.14$ , df=4, p=0.387 n.s.).

# **Supplement S1**: Physical-chemical properties of mixtures tested at the Ménudelle quarry in the Crau plain

#### Soil parameter sampling

The physical and chemical analyses of the soil were performed twice: 1) in 2016 before sowing (35 samples, 6 replicates on each of the 5 substrate mixtures + 5 on the reference grassland) and 2) in 2018 (50 samples, 5 replicates on each of the 5 substrate mixtures \* sowing/no sowing). Each 200-g sample of soil was collected by extracting the first ten centimeters of the soil layer outside the vegetation survey quadrats. Each sample was dried and sieved with a 2 mm mesh sieve. The physical and chemical analyses were performed by the soil analysis laboratory at INRA (*Institut National de la Recherche Agronomique*) in Arras, France. The reference(s) for the method used for each parameter is detailed here: pH water ratio 1:5 (Thomas 1996), CaCO<sub>3</sub> using a Bernard calcimètre (Sparks et al. 1996), organic C (Allison 1965), total N by dry combustion (Bremner 1996; Dumas 1831), C:N ratio, Cation Exchange Capacity (CEC) and exchangeable K<sup>+</sup> by Metson (Metson 1956; Ciesielski et al. 1997), available P (Olsen 1954). Particle-size distribution was studied only in 2016 in 5 fractions following the Robinson method without prior decarbonisation (clay (<0.002 mm), fine silt (0.002-0.02 mm), coarse silt (0.2-2 mm)).

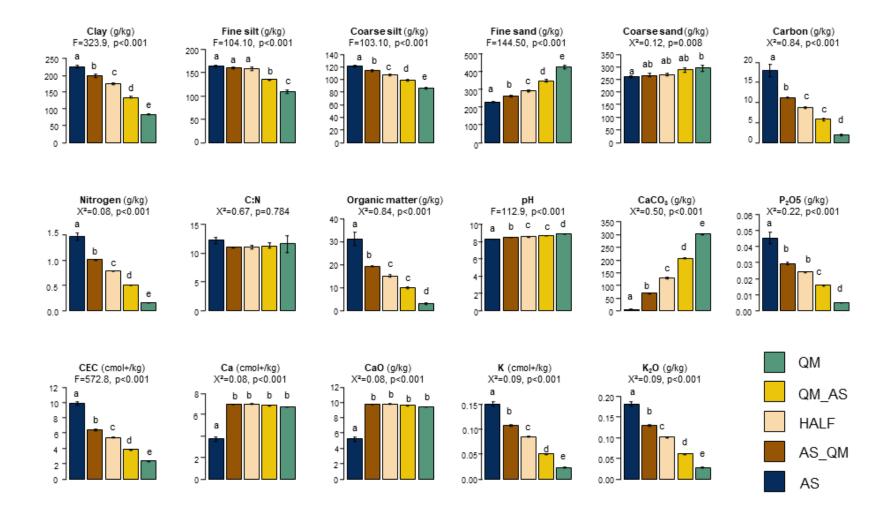
#### Data analyses of physical-chemical properties of soils

In order to compare 2016 soil physical and chemical parameter values between substrate mixtures, univariate analyses were performed. When the parametric conditions were met, an ANOVA test, followed by a post-hoc Tukey Honestly Significant Difference test, were performed whenever significant differences were encountered (Stoll 2017). When the parametric conditions were not met, Generalized Linear Models (GLM) under gamma distribution were run. These analyses were followed by pairwise contrast comparisons with a Tukey adjustment. To study the changes in the chemical parameters over years (comparing the 2016 analyses with those of 2018), a principal component analysis (Dray et al. 2020) was run on the chemical data of both years, including that of the reference grassland, to show potentially differences between years and mixtures.

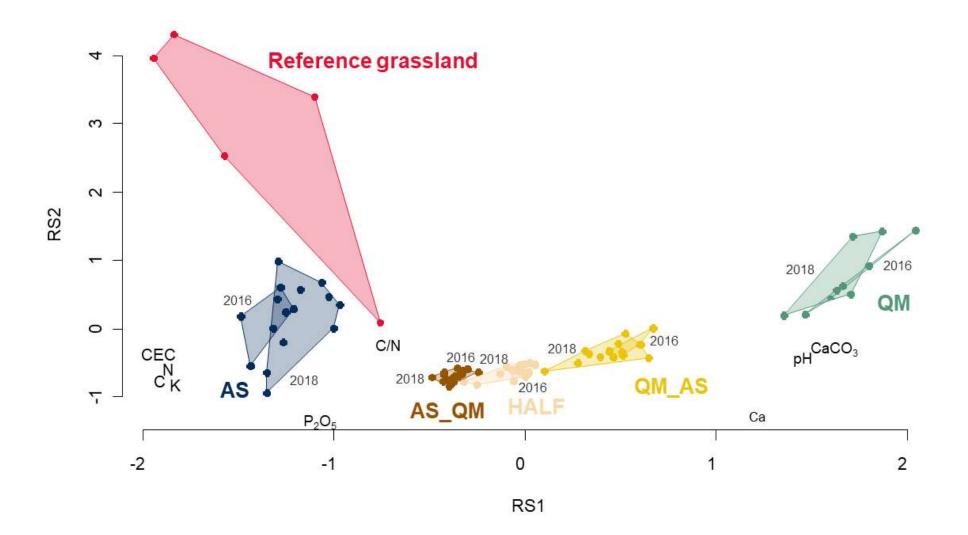
#### Patterns of change in the physical and chemical parameters of the soils

The mixture of arable soil with raw quarry material reduced the fertility of the soil (Figure S1). The concentrations of nitrogen, phosphorus and potassium were reduced by a factor of five between the substrates composed entirely of arable soil (Figure S1) and those composed exclusively of raw quarry material. The proportion of sand (fine and coarse) was much greater in the raw quarry material. In contrast, clay was found in greater quantities in the mixtures with arable soil (Figure S1). The raw quarry material was characterized by higher  $CaCO_3$  and Ca content, whereas the arable soil contained significantly more organic matter and organic carbon (Figure S1). Finally, a gradient in the Cationic Exchange Capacity was observed, with greater quantities for the arable soil compared to the raw quarry material (Figure S1).

The PCA (Figure S2) opposed the reference grassland from all experimental plots. Substrate mixtures with higher levels of arable soils had higher phosphorus, nitrogen, potassium and Cationic Exchange Capacity, while substrate mixtures with higher levels of raw quarry materials had higher  $CaCO_3$  and Ca content and higher pH.



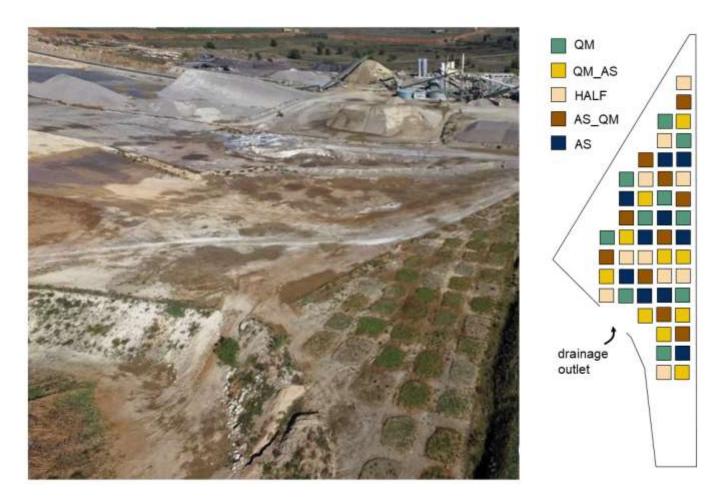
**Figure A**. The bars represent the mean ± standard error (QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil). These analyses were run on 2016 data



**Figure B**. Principal Component Analysis of chemical soil parameters on the reference grassland and on the experimental plots in 2016 and 2018. QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil

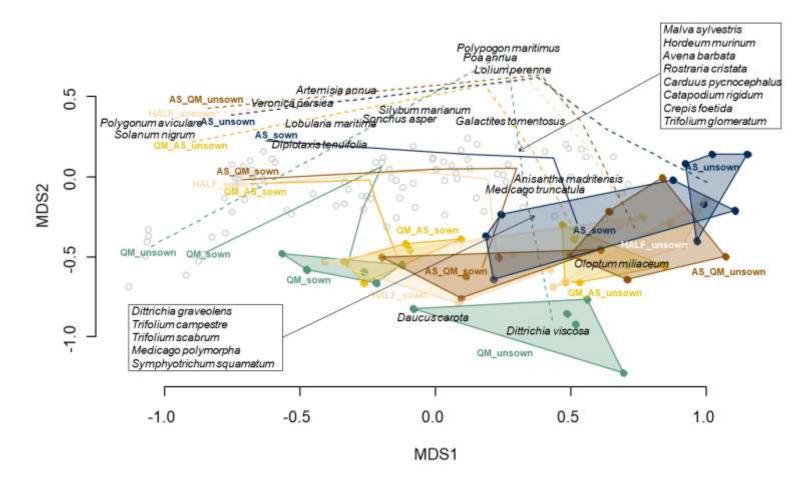
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Supplement S2. Experimental plots set up in the Ménudelle quarry in Southeastern France.

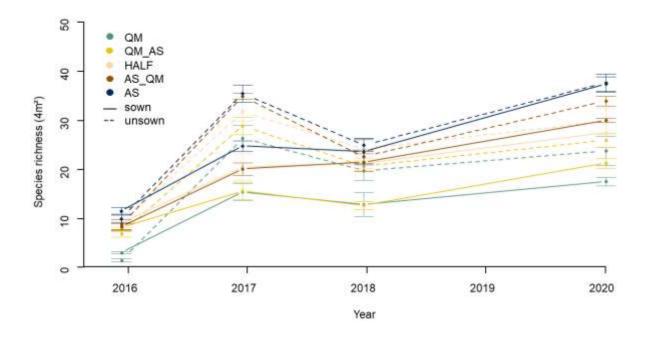


**Figure C.** Experimental plots set up in the Ménudelle quarry in 2016 (50 plots  $\times$  5 mixtures (QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil)  $\times$  2 sowing treatments + a drainage outlet) Photography: SCLM

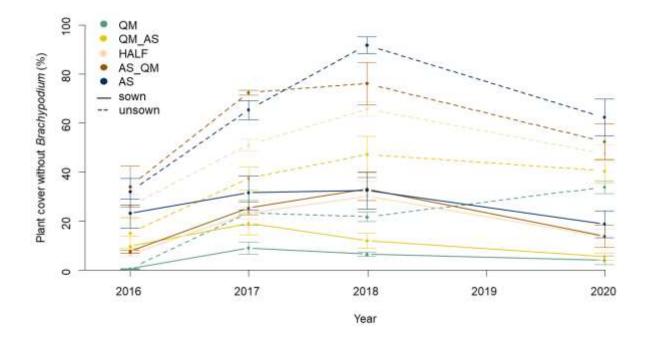
Supplement S3: Influence of substrate mixtures, Brachypodium sowing and time on plant communities



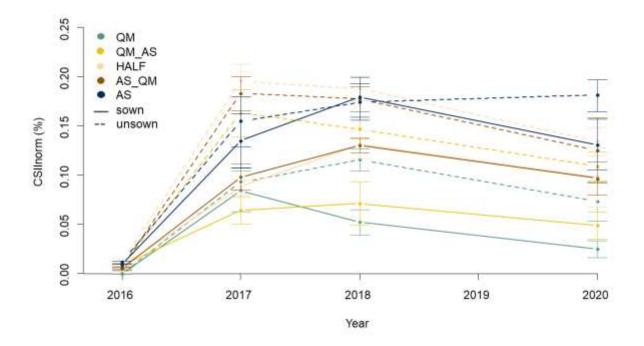
**Figure D**. Plant species composition using NMDS. Polygons indicate of plots with the same treatment in 2020 and lines the trajectory in 2016, 2017, 2018 and 2020 (NMDS stress: 0.21). Plots were sown (sown) or not (unsown) with *Brachypodium*. Substrate mixtures included QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil



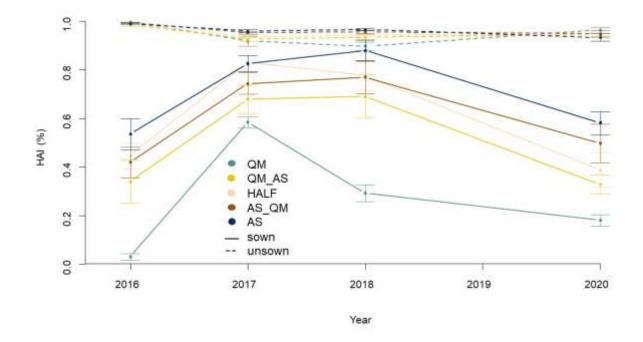
**Figure E.** Mean species richness on experimental plots in 2016, 2017, 2018 and 2020. Plots were sown (solid lines) or not (dashed lines) with *Brachypodium*. Substrate mixtures included QM – raw Quarry Material or AS – Arable Soil (see Table 1 for mixture codes). Substrate mixtures × *Brachypodium* sowing  $\chi^2$ =9.40, p=0.052; Substrate mixtures × Years  $\chi^2$ =27.49, p=0.007\*\*; *Brachypodium* sowing × Years  $\chi^2$ =34.75, p<0.001\*\*\*. QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil



**Figure F**. Mean plant cover (without *Brachypodium*) on experimental plots in 2016, 2017, 2018 and 2020. Plots were sown (solid lines) or not (dashed lines) with *Brachypodium*. Substrate mixtures included QM – raw Quarry Material or AS – Arable Soil (see Table 1 for mixture codes). Substrate mixtures × *Brachypodium* sowing  $\chi^2$ =152.13, p<0.001\*\*\*; Substrate mixtures × Years  $\chi^2$ =49.96, p<0.001\*\*\*; *Brachypodium* sowing × Years  $\chi^2$ =47.98, p<0.001\*\*\*. QM: 100% raw quarry material; QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil

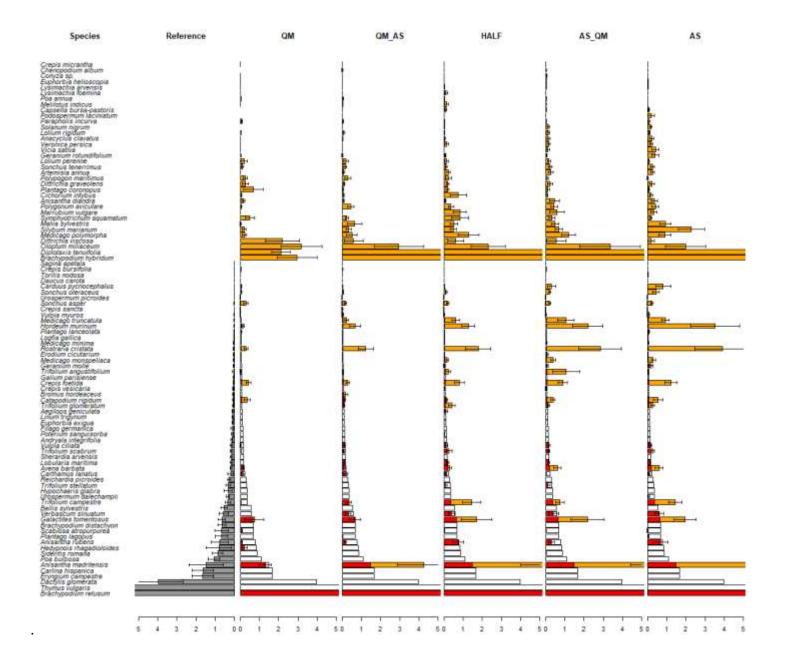


**Figure G**. Mean CSII on experimental plots in 2016, 2017, 2018 and 2020. Plots were sown (solid lines) or not (dashed lines) with *Brachypodium*. Substrate mixtures included QM – rawQuarry Material or AS – Arable Soil (see Table 1 for mixture codes). Substrate mixtures × *Brachypodium* sowing × Years  $\chi^2$ =26.24, p=0.010\*. QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil



**Figure H**. Mean HAI on experimental plots in 2016, 2017, 2018 and 2020. Plots were sown (solid lines) or not (dashed lines) with *Brachypodium*. Substrate mixtures included QM – rawQuarry Material or AS – Arable Soil (see Table 1 for mixture codes). Substrate mixtures × *Brachypodium* sowing  $\chi^2=152.13$ , p<0.001\*\*\*; Substrate mixtures × *Brachypodium* sowing × Years  $\chi^2=105.95$ , p<0.001\*\*\*. QM: 100% raw quarry material; QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil

**Supplement S4**: Structures of plant communities recorded on the different substrate mixtures tested against the reference grassland plant community (grey: mean abundance of species recorded in the reference; white: abundance lacking in relation to the reference; orange: abundance of target species present in the community assessed; red: excess abundance of species present in the communities assessed and not in the reference). (QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil; HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil)



Supplement S5: Species recorded in 2020 on the different substrate mixtures for the two sowing treatments (with or without sowing of *Brachypodium*) and the reference grassland. QM: 100% of raw Quarry Material; QM\_AS: 75% of raw Quarry Material and 25% of Arable Soil;
HALF: 50% of each; AS\_QM: 75% of Arable Soil and 25% of raw Quarry Material; AS: 100% of Arable Soil

		QM		QI	M_AS	HALF		AS_QM		AS		Grassland
Genus	species	sown	unsown	sown	unsown	sown	unsown	sown	unsown	sown	unsown	Grassianu
Aegilops	geniculata				х	х	х	х	х	х	х	х
Aira	cupaniana									х		х
Ajuga	iva					х						
Anacyclus	clavatus	х			х	х	х	х	х	х	х	
Andryala	integrifolia					х		х		х	x	х
Anisantha	diandra	х	x		x							
Anisantha	madritensis	х	x	х	х	х	x	х	x	х	x	х
Anisantha	rubens	х	x	х	х	х	x	х	х	х	x	х
Arenaria	leptoclados					х			х			х
Artemisia	annua						x	х	х	х	x	
Artemisia	verlotiorum						х				х	
Asphodelus	ayardii											х
Avena	barbata		х	х	х	х	х	х	х	х	х	х
Bellis	sylvestris											х
Bituminaria	bituminosa		x				x					
Bothriochloa	ischaemum											х
Brachypodium	distachyon											х
Brachypodium	hybridum	х	х	х	х	х	x	х	х	х	x	
Brachypodium	retusum	х	x	х	х	х		х		х		х
Bromus	hordeaceus	х			х	х	х			х	х	х
Bupleurum	semicompositum						х	х			х	
Carduus	nigrescens				х							х
Carduus	pycnocephalus	х			х	х	х	х	х	х	х	х
Carduus	tenuiflorus						х			х		

Carlina	hispanica											x
Carlina	lanata						х	х	х		х	х
Carthamus	lanatus		х	х	х		х	х	х	х	х	
Catapodium	rigidum	х	х	х	х	х	х	х	х	х	х	х
Centaurium	erythraea											х
Centaurea	melitensis											х
Cerastium	glomeratum	х									х	
Cerastium	pumilum		х				х		x			х
Chondrilla	juncea		х				х					
Cichorium	intybus					х					х	
Clinopodium	nepeta		х									
Convolvulus	arvensis						х	х	х	х		
Convolvulus	cantabrica			х	х				х	х	х	
Conyza	sp.		х		х			х	х	х	х	
Crepis	bursifolia		х	х	х			х	x	х		
Crepis	foetida	х	х	х	х	х	х	х	х	х	х	х
Crepis	micrantha		х	х		х	х	х		х		
Crepis	sancta			х	х	х		х	x	х	х	х
Crepis	vesicaria		х	х		х	х	х	х	х	х	х
Cynodon	dactylon	х				х				х	х	
Cynosurus	echinatus						х					х
Dactylis	glomerata				х	х	х		x			х
Daucus	carota	х	х	х	х	х	х	х	x	х		
Diplotaxis	tenuifolia	х	х	х	х	х	x		x	х	х	
Dittrichia	graveolens	х	х	х	х	х	х	х	х	х	х	
Dittrichia	viscosa	х	х	х	х	х	х	х	х	х	х	
Echium	vulgare		х		х		х		х		х	
Erodium	cicutarium							х	х		х	х
Erodium	malacoides						х		х	х	х	

Erodium	moschatum	х		х		х			x			
Eryngium	campestre						х					х
Euphorbia	cyparissias					х						x
Euphorbia	exigua											x
Euphorbia	falcata											x
Euphorbia	helioscopia		х						х	x	х	
Euphorbia	segetalis								х			
Euphorbia	serrata		х									
Filago	germanica		х	х	х	х	х			x	х	х
Filago	рудтаеа		х							х		х
Filago	pyramidata		х									
Foeniculum	vulgare		х		х		х	х				
Galactites	tomentosus	х	х	х	х	х	х	х	х	х	х	х
Galium	murale											х
Galium	parisiense	х			х				х			х
Geranium	molle			х	х	х	х			x	х	х
Geranium	rotundifolium	х	х			х	x		х	x	х	
Hedypnois	rhagadioloides		х					х		x		х
Hippocrepis	ciliata							х				
Hirschfeldia	incana					х		х		x		
Hordeum	murinum	х	х	х	х	х	х	х	х	x	х	
Hypericum	perfoliatum						x		х			
Hypochaeris	glabra					х			х	x	х	х
Lactuca	serriola										х	
Lagurus	ovatus										х	
Lathyrus	cicera									х		
Leontodon	tuberosus							х				
Limonium	echioides							х				
Linaria	arvensis											x

Linum	strictum								x		х	x
Linum	trigynum											х
Lobularia	maritima					х	х		х	х	х	х
Logfia	gallica									х		х
Lolium	rigidum	х	х	х	х	х	х	х	х	х	х	
Lysimachia	arvensis	х			х		х	х	x	х	х	
Lysimachia	foemina		х		х		х		х			
Lysimachia	linum-stellatum											х
Malva	sylvestris	х			х	х	х		x	х	х	
Marrubium	vulgare					х	х		х	х	х	
Medicago	minima	х		х		х		х		х	х	х
Medicago	monspeliaca	х	х	х	х	х	х	х	x	х	х	х
Medicago	orbicularis	х		х		х		х				
Medicago	polymorpha	х	х	х	х	х	х	х	х	х	х	
Medicago	rigidula	х			х	х	х	х			х	х
Medicago	truncatula	х		х	х	х	х	х	х	х	х	х
Melilotus	indicus	х		х	х	х		х	х	х		
Misopates	orontium			х						х		
Neatostema	apulum		х									х
Oloptum	miliaceum	х	х		х		х		х		х	
Pallenis	spinosa			х	х	х		х	х			
Papaver	rhoeas						х				х	
Parapholis	incurva	х	х	x	х	х	х	х	x		х	
Petrorhagia	prolifera							х				х
Plantago	afra											х
Plantago	bellardii					х						х
Plantago	coronopus	х	х		х	х	х	х	х	х	х	
Plantago	lagopus	х			х			х		х		х
Plantago	lanceolata	x	х			х					х	

Роа	bulbosa											x
Podospermum	laciniatum				х	х	х	х	х	х	х	
Polygonum	aviculare							х				
Polypogon	maritimus	х										
Polycarpon	tetraphyllum											х
Poterium	sanguisorba		х	х		х	х					х
Prospero	autumnale											х
Psilurus	incurvus											х
Reichardia	picroides											х
Rostraria	cristata	х	х	х	х	х	х	х	х	х	х	х
Rumex	pulcher								х			х
Sagina	apetala		х		х		х		х	х	х	х
Scabiosa	atropurpurea	x	х	х	х			х			х	х
Scolymus	hispanicus									х	х	
Scorpiurus	muricatus		х							х		
Senecio	vulgaris											х
Sherardia	arvensis							х	х	х	х	х
Sideritis	romana		х						х			х
Silene	gallica											х
Silene	nocturna			x			х	х	х	х		х
Silybum	marianum	х	х	х	х	х	х	х	х	х	х	
Sonchus	oleraceus				х				х			х
Sonchus	asper	x	х	х	х	х	х	х	х	x	х	х
Sonchus	tenerrimus					х		х		х	х	
Stipa	capillata											х
Symphyotrichum	squamatum	х	х	х	х	х	х	х	х	х	х	
Taeniatherum	caput-medusae											х
Thymus	vulgaris											х
Tolpis	umbellata											х

Torilis	nodosa								х			
Tragopogon	porrifolius		х			х		х	х			
Trifolium	angustifolium	х	х	х	х	х	х	х	х	х	х	
Trifolium	campestre	x	х	х	х	х	х	х	x	x	x	х
Trifolium	cherleri							х				х
Trifolium	glomeratum			х	х	х	х	х	х	х	x	х
Trifolium	scabrum	x	х	х	х	х	х	х	х	х	х	х
Trifolium	stellatum					х		х	х			х
Trifolium	suffocatum						х				x	
Trifolium	tomentosum						х			х		
Urospermum	dalechampii		х									х
Urospermum	picroides		х		х	х	х	х	х	х	х	х
Valantia	muralis								х			
Verbascum	sinuatum		х	х	х		х		х	х	х	х
Veronica	arvensis							х		х	х	
Veronica	persica				x		х	х	х		х	
Veronica	polita										х	
Vicia	lathyroides									x		
Vicia	sativa						х	х		х	x	
Vulpia	ciliata	x	х	x	х	х	х	х	х	x	x	х
Vulpia	myuros	х	х	х	х	х	х		х	х	х	х