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More warm-adapted species in soil seed banks than in herb layer plant communities across Europe

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1 **RESEARCH ARTICLE**

More warm-adapted species in soil seed banks than in herb layer plant communities across Europe

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54

55 Abstract

Responses to climate change have often been found to lag behind the rate of warming that has
 occurred. In addition to dispersal limitation potentially restricting spread at leading range margins,
 the persistence of species in new and unsuitable conditions is thought to be responsible for apparent
 time lags.

60 2. Soil seed banks can allow plant communities to temporarily buffer unsuitable environmental conditions, but their potential to slow responses to long-term climate change is largely unknown. As 61 local forest cover can also buffer the effects of a warming climate, it is important to understand how 62 63 seed banks might interact with land cover to mediate community responses to climate change. 3. We first related species-level seed bank persistence and distribution-derived climatic niches for 64 840 plant species. We then used a database of plant community data from grasslands, forests and 65 intermediate successional habitats from across Europe to investigate relationships between seed 66 banks and their corresponding herb layers in 2763 plots in the context of climate and land cover. 67 68 **4.** We found that species from warmer climates and with broader distributions are more likely to have a higher seed bank persistence, resulting in seed banks that are composed of species with 69 warmer and broader climatic distributions than their corresponding herb layers. This was consistent 70 across our climatic extent, with larger differences (seed banks from even warmer climates relative 71 to vegetation) found in grasslands. 72

5. Synthesis. Seed banks have been shown to buffer plant communities through periods of
environmental variability, and in a period of climate change might be expected to contain species
reflecting past, cooler conditions. Here, we show that persistent seed banks often contain species
with relatively warm climatic niches and those with wide climatic ranges. Although these patterns
may not be primarily driven by species' climatic adaptations, the prominence of such species in

78	seed banks might still facilitate climate-driven community shifts. Additionally, seed banks may also
79	be related to ongoing trends in the spread of widespread generalist species into natural habitats,
80	while cool-associated species may be at risk from both short- and long-term climatic variability and
81	change.
82	
83	Keywords: Climate change, Climatic debt, Dispersal, Plants, Seed longevity, Seedbank,
84	Thermophilisation
85	
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87	Introduction
88	Ongoing changes in the climate system are having profound effects on the Earth's ecosystems.
89	Species are shifting their ranges poleward and to higher altitudes (Chen et al., 2011; Kelly &
90	Goulden, 2008), the timings of life-history events are shifting (Parmesan & Yohe, 2003; Wolkovich
91	et al., 2012) and populations are going locally extinct (Wiens, 2016). Changes in individual and
92	population-level responses are in turn reflected in changes at the community level, with species that
93	are associated with warmer climates increasing at the expense of those species associated with
94	cooler climates (Auffret & Thomas, 2019; De Frenne et al., 2013; Devictor et al., 2012; Fadrique et
95	al., 2018). However, the responses that species and communities exhibit following climate change
96	do not always follow this expected pattern, or are slower than would be predicted from the
97	magnitude of warming that has occurred (Ash et al., 2017; Becker-Scarpitta et al., 2019; Bertrand et
98	al., 2011).

99

The rate at which plant species respond to climate change can be driven both by their ability to
colonise new areas, as well as by the extent that populations are able to persist after local conditions
have become climatically unsuitable (Alexander et al., 2018). Both of these responses can be related
to the species' potential to disperse. Studies of changes in plant communities over time have shown

that non-native species and other cosmopolitan generalists with large or expanding ranges have
increased their distributions during the past decades (Auffret & Thomas, 2019; Finderup Nielsen et
al., 2019; Staude et al., 2020), presumably via effective spatial dispersal. Nonetheless, the rate at
which the climate is changing means that even species that are able to disperse long distances may
still be limited when it comes to the possibility to effectively track their shifting climatic niche
(Alexander et al., 2018; González-Varo et al., 2021).

110

While dispersal limitation in space might restrict some species' ability to effectively expand their 111 distributions in a warming climate, dispersal through time in seed banks could potentially contribute 112 to local persistence of cold-adapted populations, thus explaining observed slow responses that have 113 been exhibited at the trailing end of species' distributions and at the community level. By buffering 114 115 short and long-term environmental variability, seed banks are considered to play an important role in plant population and community dynamics (DeMalach et al., 2021; Eriksson, 1996). As such, 116 species that form persistent seed banks may have a reduced risk of local extinction following 117 environmental change (Auffret et al., 2017), with seed bank communities often reflecting historical 118 patterns of land use and management (Karlík & Poschlod, 2014; Plue et al., 2008). Therefore, seed 119 banks might also be expected to reflect historical climatic conditions, and evidence does suggest 120 that they may contribute to the maintenance of species diversity in the face of climate and land-use 121 change (Plue et al., 2021; Vandvik & Goldberg, 2006). However, experimental studies have shown 122 that seed bank size and richness can be directly negatively affected by changing climatic conditions 123 (Basto et al., 2018; Eskelinen et al., 2021). It is clearly important to improve our understanding of 124 the role of seed banks in community responses to climate change. However, it should also be noted 125 126 that despite evidence that seed banks can buffer communities in periods of environmental uncertainty, the species found in a seed bank are not a random subset of the species found in the 127 herb layer. Instead, they represent a group of species whose persistent seeds are part of a specific 128

- 129 life strategy, and which might also differ from many species in the herb layer in terms of other
- 130 functional traits (Pakeman & Eastwood, 2013; Thompson et al., 1998).
- 131

132 The rate at which communities respond to warming is not only determined by species characteristics, but also by the extent to which they are exposed to climate change. Plant 133 communities in different habitats are often exposed to different microclimates, even if they are in 134 close proximity. For example, the buffering effect of forest canopies results in cooler temperatures 135 in summer and milder temperatures in winter, compared to open sites (De Frenne et al., 2019; 136 Morecroft et al., 1998). This is important, because the microclimates that plant communities 137 experience are key determinants in both the facilitation of positive responses to climate warming, 138 but can also reduce climate-related extirpations that might be expected from the macroclimatic 139 140 changes manifesting at larger spatial scales (Suggitt et al., 2018; Zellweger et al., 2020). Indeed, changes in forest cover at the local and landscape-level have been shown to influence rates of 141 community change over time through their effects on microclimatic conditions. The cooling and 142 stabilising effect of increased tree canopy or forest cover has been shown to reduce both 143 establishment and extinction of warmer- and cooler-adapted species, respectively (Auffret & 144 Thomas, 2019; De Frenne et al., 2013). Local forest cover can also affect relationships between soil 145 seed banks and the herb layer through the effects of microclimate (Gasperini et al., 2021), and as 146 such, land cover can be an important consideration when investigating the effects of climate change 147 148 on plant communities.

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Here we assess the relationship between seed bank persistence and climate, and in doing so explore the role that seed bank persistence in the soil may have on community responses to climate change. We consider the following key questions: [1] We first ask if a species' seed bank persistence is related to its climatic niche (calculated as the average temperature from across a species' geographic range)? [2] We then use a dataset of seed bank and corresponding herb layer community 155 plots across a climate and land-cover gradient to calculate community temperature indices - the average species temperature index within a community – to ask whether seed bank communities 156 consist of relatively cool-associated species compared to the herb layer, reflecting past, potentially 157 158 cooler communities and slowing down community responses to climate change? Finally, [3] we ask whether differences in community temperature indices between the seed bank and the herb layer are 159 related to land cover and macroclimate. Throughout our analyses, we consider the potential effects 160 of additional plant functional traits in driving observed patterns relating to species and community 161 climatic indices in the seed bank and herb layer. 162

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165 Methods

166 Data preparation

167 Seed bank database

We used an existing database of 2796 paired seed bank and herb layer presence-absence community 168 plots from across northern Europe, including data from southern France to mid-Sweden and 169 Norway, and from western England in the west to Estonia in the east (Figure S1 and Table S1 in the 170 Supplementary Information). The database consists of presence-absence occurrences from 54 171 original data sets, and as such reflects a range of sampling techniques. Nonetheless, all seed bank 172 communities were sampled through the collection of soil and subsequent greenhouse germination 173 assays with the aim of quantifying the long-term persistent seed bank, while the herb layer was 174 assessed in relevés from the same sampling locations, with each individual study designed to 175 compare communities in the seed bank and herb layer. Seed bank plot sizes ranged from 0.0015 -176 0.62 m^2 area and 0.03 - 0.2 m depth, while time of sampling also varied. Vegetation relevés ranged 177 from 1 to 400 m². The data cover a broad temperature gradient, and the database includes mean 178 179 annual temperature for each plot at the approximate time of sampling extracted from the CHELSA time-series database v1.2, 1978-2014 (Karger et al., 2017, 2018). Each plot was also assigned to 180

181 one of three broad land cover categories: low-intensity managed grassland, mature forest, and intermediate successional habitats, which included abandoned grasslands with shrubs and young 182 forests, such as post-agricultural forests. These categories were spread across the climate gradient, 183 184 with seven of the nine countries covered by the database containing plots from all three categories. Community data were collected between 1978 and 2014, with the majority of datasets (67%, 185 including 85% of all plots) collected since 2000, by which time European surface temperatures had 186 already warmed by almost 1 °C compared to pre-industrial estimates (European Environment 187 Agency, 2022). A previous study using the database found that species richness was higher in the 188 seed bank than the herb layer, and that the seed bank was less affected by climate and land use 189 variables than the herb layer, indicating a potential buffering effect (Plue et al., 2021). 190

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192 Species' climatic niches

Species' climatic niches were taken from the ClimPlant database (Vangansbeke et al., 2021a, 193 2021b), which estimates the realised climatic niches of 968 forest understorey species based on the 194 climatic conditions from their European range (bordered by the Atlantic ocean, Arctic ocean, Ural 195 mountains and Sahara desert) averaged across the 1970-2000 reference period. Briefly, scanned 196 European range maps for each species were georeferenced and digitised using a geographic 197 information system. These digitisations were then overlain with the WorldClim dataset (Fick & 198 Hijmans, 2017), and temperature and precipitation data from each species were extracted by 199 randomly sampling one thousand 20×20 km grid squares (with replacement) from within that 200 species' range. Because our seed bank database also included species from open and intermediate 201 successional habitats, we used the same methodology to complement the ClimPlant database with 202 203 climatic niches for 200 additional species (i.e., all species in the seed bank database that were missing from ClimPlant and for which we could find European range maps). This way, we obtained 204 climatic niches for 93% of species from the seed bank database (including species only observed in 205 the herb layer). For this study, we extracted two temperature indices for all possible species. First, 206

207 the mean annual temperature within the species' distribution range (hereafter species mean 208 temperature index, where higher values indicate a warmer, more southerly distribution), and second the range of the mean annual temperature within the species' distribution range, excluding the 5% 209 210 warmest and 5% coldest mean temperatures to avoid the influence of extreme values (hereafter species temperature range index, where higher values indicate a more widespread climatic 211 distribution. Across the ClimPlant data set, there was little evidence of correlation between species 212 mean temperature index and species temperature range index. Although the metrics are somewhat 213 related (the most widespread species would have a species temperature index approximating the 214 215 European mean average temperature), the existence of (for example) narrow-ranged species in both more northerly and southerly regions contributes to there being only a very slight negative 216 correlation between the two indices (Pearson correlation coefficient -0.1; Figure S2). We therefore 217 218 consider the indices to be adequately independent for analysis purposes. When considering only species that were present in the seed bank database, the correlation coefficient was -0.089. 219

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221 Plant functional traits

The LEDA traitbase (Klever et al., 2008) contains information on seed bank persistence for 1586 222 European plant species. We calculated the seed longevity index (Thompson et al., 1997), which is 223 the proportion of records reporting a species to have a persistent, compared to a transient seed bank, 224 based on naturally-buried seeds. As such, the metric reflects a combination of seed physiology and 225 environmental conditions that can affect whether a seed persists in the soil. Here, we calculated for 226 each species the proportion of rows in the LEDA traitbase for which it is listed as being 'long-term 227 persistent' (at least five years) or 'short-term persistent' (between one and five years), as opposed to 228 229 'transient' (less than one year). Rows in which seed bank status was 'present' were removed, because it was not possible to tell whether the seed bank was persistent or transient. The resulting 230 index therefore ranges from 0 (never recorded as having a persistent seed bank) to 1 (always 231

recorded as having a persistent seed bank).

233	Three additional plant traits that have been associated with seed bank persistence were also
234	extracted from the LEDA traitbase: [1] plant lifespan (perennial or annual, including biennial),
235	whereby annual species are more often found in seed banks (Gioria et al., 2020; Thompson et al.,
236	1998); [2] seed mass (mg), whereby smaller seeds are more likely to have higher seed bank
237	persistence (Gioria et al., 2020; Hodkinson et al., 1998); and [3] seed number ('per ramet, tussock
238	or individual plant'), whereby species producing more numerous seeds are logically more likely to
239	be detected in seed banks, while there is an apparent trade-off between seed size and seed
240	production (Leishman, 2001). Values for each species were calculated as the geometric mean of all
241	available values for that species, to reduce the effect of extreme values.
242	

243 Data analysis

244 Seed bank persistence and species temperature indices (Question 1)

As a first step to understand community-level climate associations across the seed bank and herb 245 layer, we assessed the relationship between a species' seed bank persistence and its climatic niche. 246 Therefore, we created two binomial generalised linear models (function: glm, family: quasibinomial 247 due to overdispersion) in the R statistical environment (version 4.2.0; R Core Team, 2022), where 248 the response variable was the seed bank longevity index calculated above. In the first model (Model 249 1a), the predictor variable was the species mean temperature index, and in the second model (Model 250 1b), the predictor variable was the *species temperature range index*. Because our response variable 251 (seed bank longevity index) is a proportion of successes/failures (i.e. observations of long-term 252 persistence or otherwise), the number of 'trials' needed to be included as a weight in the binomial 253 models. For this, we used the number of seed bank assessments (i.e. data rows) for each species in 254 255 the LEDA traitbase. To maximise the power of this analysis, we included all 840 species for which both species temperature indices and seed bank longevity index were available, even considering 256 species that were not present in the seed bank database. Statistical significance of predictor 257 variables in generalised linear models was determined using a p-value threshold of 0.05. 258

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260 To assess the potential correlates of other plant traits on seed longevity index, we re-ran the above models with seed bank longevity index as the response variable and either species mean 261 262 temperature index or species temperature range index as the predictor variable, this time including plant lifespan, seed mass and seed number as additional predictor variables (Models 1c-d). 263 Correlations between species' climatic niche values and the three additional traits identified no 264 strong collinearity (Table S2), with the largest Pearson correlation coefficients being 0.15 between 265 species mean temperature index and seed mass, and -0.15 between species mean temperature index 266 267 and plant lifespan (converted to binary, with annuals as 0 and perennials as 1). The order of predictors in the model formulae did not affect the model outputs. 268 269 270 Comparison of community temperature indices between the seed bank and herb layer (Question 2) In the next step, we calculated two community temperature indices for each seed bank and herb 271 layer community for each plot in the seed bank database. First, we calculated the community mean 272 temperature index (CMTI; often referred to in the literature as the community temperature index) as 273 the average of the *species mean temperature index* for all species present within the seed bank or 274 275 herb layer community. Second, we calculated a *community temperature range index* (CTRI) as the mean of every species temperature range index within a community, in which higher values 276 indicate on average a higher relative number of climatically widespread species in the community. 277 278 Community indices were not weighted by abundance because the seed bank database only contains presence-absence information. Although abundance-weighted community climate indices are useful 279 in detecting more subtle shifts in climate-driven community composition (Lindström et al., 2013), 280 281 the use of presence-absence data is also common practice in studies where abundance data are not available, detecting both shifts in plant communities and their environmental drivers (Auffret & 282 Thomas, 2019; Lenoir et al., 2013). Differences in sampling area between the seed bank and herb 283 layer can affect estimations of community similarity (Plue et al., 2021), but here we assume that the 284

recording of species with particular climatic niche values are not related to the area of ground sampled. Thirty-three plots had seed bank or herb layer communities consisting only of species for which temperature indices could not be calculated, and were therefore excluded from the community-level analyses, leaving 2763 plots (Grassland: 1298, Intermediate: 864, Forest: 601).

Next, the community mean temperature index was regressed against the gridded (CHELSA) 290 macroclimate mean annual temperature data for each plot to verify if a climatic control on seed 291 bank and herb layer communities was present in the dataset. This was performed using a linear 292 mixed effects model (function: lmer in R package lme4 version 1.1.29; Bates et al., 2014). Data 293 were arranged so that each community occupied a separate row, that is, there were two data points 294 per plot, one for seed bank and one for herb layer. The response variable in the model (Model 2a) 295 296 was *community mean temperature index*. Fixed predictor variables were annual mean temperature (at the plot) and source (seed bank or herb layer), as well as their interaction. This allowed us to 297 first test for climatic control of the plant communities, but also whether there was a difference in 298 community mean temperature index between the seed bank and herb layer, and whether climatic 299 control of the community differed according to source. Plot identity, nested into the identity of the 300 original dataset from which the plot was taken (of the 54 component datasets) was added as a 301 random effect. Another model (Model 2b) with *community temperature range index* as the 302 response, source as the predictor, and the same random effect structure as the previous model tested 303 whether seed bank and herb layer communities differed in terms of community temperature range 304 index. Significance of mixed model effects was estimated using bootstrapped 95% confidence 305 intervals (CIs; R function: confint), with significant effects defined as CIs not including zero. 306 307

To test the potential effects of other plant traits driving temperature associations at the community level, we calculated mean values for each of the three additional traits (plant lifespan, seed mass, seed number) for the species present in the seed bank and herb layer in each plot. Correlation 311 analyses showed no strong collinearity in average trait values across communities (Table S4). We 312 then re-ran our two linear mixed models designed to assess differences in community thermal indices across the seed bank and herb layer, with community mean temperature index and 313 314 community temperature range index as response variables. Fixed predictor variables were source (seed bank or herb layer), as well as the community mean values of each additional trait. Plot 315 identity, nested into original dataset identity was included as a random effect (Models 2c and 2d). 316 Mean annual temperature at the plot was not included in the community mean temperature index 317 model, because we were not interested in climatic control of the communities in these additional 318 319 analyses.

320

321 Environmental drivers of differences in community temperature indices (Question 3)

322 To explore the potential drivers of any differences in community temperature index between the seed bank and herb layer, we built a further linear mixed model (Model 3a). The difference between 323 seed bank and herb layer *community mean temperature index* was the response variable (seed bank 324 index minus herb layer index, with positive values indicating that the seed bank is characterised by 325 warm-adapted species with higher species mean temperature indices relative to the herb layer). 326 Predictor variables were land-use category (categorical variable with intermediate successional 327 habitats as base factor for comparison) and mean annual temperature from the seed bank database. 328 the seed bank *community temperature range index* and the first two eigenvectors of a principal 329 coordinates analysis derived from a neighbour matrix (PCNM) of the spatial coordinates of each 330 plot (function: pcnm in R package vegan version 2.6.2; Borcard & Legendre, 2002; Oksanen et al., 331 2016). These variables were included to account for spatial autocorrelation because the nested 332 333 random effect structure above was not possible because there was only one row per plot in the data frame. The original dataset of each plot was included as a single random effect. Two-way 334 interactions were included for predictor variables, excluding PCNM eigenvectors. In this model, 335

- numerical predictor variables were standardised (mean = 0 and sd = 1; R function: scale) to allow
- interpretation of both main effects and interactions (Schielzeth, 2010).
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- 339

340 **Results**

- 341 Seed bank persistence and temperature indices (Question 1)
- Comparing species' seed bank longevity indices with their temperature indices revealed that species with warmer climatic niches (Model 1a) and those that are climatically widespread (Model 1b), are more likely to have a higher seed bank longevity index (*species mean temperature index*:
- Regression coefficient = 0.12, p < 0.001; species temperature range index: Regression coefficient =
- 0.11; p < 0.001; Figure 1, Table S3). Including plant lifespan, seed mass and seed number in the
- 347 models together with the temperature indices resulted in no significant effects on seed bank
- longevity index, indicating that when holding other traits constant, there are no clear associations
- 349 between each trait and seed bank persistence (Models 1c-d; Table S3). Separate models with only
- 350 one of the additional traits as a predictor variable confirmed previous findings that annual species
- are more likely to have persistent seeds, while there was no clear effect of seed mass or seed size
- 352 (Models 1e-g; Table S3).
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363 *Figure 1.* Relationships between seed bank longevity (where 0 = species always has transient seed banks, and 1 = species always has persistent seed banks, calculated from the LEDA traitbase) and 364 a) species mean temperature index (mean annual temperature across the species' range), and b) 365 species temperature range index (90% range of annual temperature from across the species' 366 range). In each panel, grey points are individual species and the red line is the modelled 367 relationship with bootstrapped 95% confidence intervals calculated from Model 1a and 1b. Note 368 that the modelled relationship is from a binomial logistic regression where the trend line indicates 369 for each value of X the probability of Y being one. 370

371

372

373 *Comparison of community temperature indices (Question 2)*

Both seed bank and herb layer communities appear to be subject to macroclimatic control, with the

- 375 *community mean temperature index* being higher in plots that have higher mean annual
- temperatures (Model 2a, Regression coefficient = 0.2, CI = 0.16-0.24; Figure 2). The level of
- 377 climatic control did not vary between the seed bank and herb layer, as evidenced by the non-
- 378 significant interaction between mean annual temperature and source (Model 2a, Regression

379	coefficient = 0.003 , CI = $-0.01-0.01$). Contrary to our expectations, seed bank communities
380	contained on average species that have warmer climatic ranges compared to the herb layer (Model
381	2a, Regression coefficient = 0.23 , CI = $0.13-0.36$, Figure 3a), as well as species that were
382	climatically more widespread (Model 2b, Regression coefficient = 0.64, CI = 0.59-0.68; Figure 3b).
383	Models including community mean values of plant lifespan, seed mass and seed number showed
384	that despite significant effects on both community mean temperature index (Model 2c; higher
385	values in communities containing relatively more annuals, more large-seeded species, and species
386	producing more seeds) and community temperature range index (Model 2d; higher values in
387	communities containing relatively more annuals, more large-seeded species and species producing
388	fewer seeds), the pattern that seed banks contained relatively warmer and more widespread species
389	remained significant (Table S5).
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Figure 2. Relationships between the local macroclimate mean annual temperature and the
community mean temperature index (CMTI) of (a) the herb layer and (b) the seed bank in 2763
plots across Europe. In each panel, points are individual plots and the line is the modelled
relationship with 95% confidence intervals, calculated using the visreg package from the outputs of
Model 2a (version 2.7; Breheny & Burchett, 2017).

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414 Environmental drivers of differences in thermal indices (Question 3)

The difference in community mean temperature index between the seed bank and herb layer was

- independent of the mean annual temperature, but was higher in grasslands compared to intermediate
- 417 successional habitats (Table 1). Communities having a relatively higher representation of
- climatically widespread species were also more likely to have higher *community mean temperature*
- 419 *index* in the seed bank than the herb layer, although this effect was smaller in plots with relatively
- 420 higher mean annual temperatures (negative interaction between temperature and seed bank

421 community temperature range index). Significant negative interactions were also found between 422 community temperature range index in the seed bank and forest and grassland habitat category. In 423 other words, the effect of larger numbers of climatically widespread species in the seed bank that 424 are potentially driving the pattern of warmer seed bank communities and cooler herb layer 425 communities, was larger both in forest and grassland plots compared to intermediate successional 426 habitats.

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Figure 3. Differences in: a) community mean temperature index (CMTI); and b) community temperature range index (CTRI) according to habitat category, calculated as the index of the seed bank community minus the index of the corresponding herb layer community (jittered coloured points). Boxes show median and interquartile range, with whiskers indicating range excluding outliers. Notches represent 95% intervals around the median. White points indicate the mean values.

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Table 1. Linear mixed model (Model 3a) outputs showing the effect of temperature, land use and community temperature range index of the seed bank on the difference between the community temperature index of seed bank and the herb layer. Arrow symbols represent significant positive (i.e. higher values of the predictor result in relatively warmer seed bank communities), negative or non-significant effects, as defined by whether 95% confidence intervals include zero.

		Coefficient	Standard Error	t value	CI Lower 2.5%	CI Upper 97.5%
Main effects						
Mean annual temperature (MAT)	\leftrightarrow	-0.047	0.083	-0.57	-0.20	0.11
Grassland habitat (compared to intermediate successional)	1	0.1	0.046	2.06	0.002	0.18
Forest habitat (compared to intermediate successional)	\leftrightarrow	-0.051	0.11	-0.45	-0.30	0.18
Seed bank community temperature range index (SB - CTRI)	1	0.15	0.026	5.91	0.10	0.20
Interaction effects						
MAT : Grassland	\leftrightarrow	0.11	0.07	1.55	-0.03	0.25
MAT : Forest	\leftrightarrow	-0.015	0.10	-0.15	-0.23	0.20
MAT : SB - CTRI	\downarrow	-0.1	0.021	-4.77	-0.14	-0.06
SB – CTRI : Grassland	\downarrow	-0.26	0.036	-7.05	-0.33	-0.19
SB – CTRI : Forest	\downarrow	-0.12	0.049	-2.4	-0.21	-0.02
Spatial autocorrelation						
PCNM 1	\leftrightarrow	-0.022	0.09	-0.24	-0.20	0.16
PCNM 2	1	0.13	0.05	2.46	0.03	0.24

448 **Discussion**

Our study of the community temperature indices of 2763 European seed bank and herb layer plots showed – surprisingly – that seed banks are associated with more warm-associated species than their corresponding aboveground herb layer communities. Despite the eco-evolutionary role of seed banks buffering periods of environmental instability to allow population re-establishment, our results indicate that in a period of rapid climate change, rather than lagging behind changing plant communities, seed banks may instead have the potential to give plant communities a head start on the changes to come.

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Why are species with relatively high seed bank persistence associated with warmer climates and 457 large climatic ranges? Studies indicate that increased temperatures are likely to diminish the ability 458 for species to persist in the soil for long periods due to increased seed damage (Ooi et al., 2009), 459 and that higher temperatures and increased drought frequencies are likely to directly or indirectly 460 impact seed banks negatively (Walck et al., 2011). However, it does not necessarily follow that 461 plant species with a warmer climatic niche have lower seed bank persistence. In Europe, southerly 462 regions with warmer climates are also those that experience more extreme events in terms of heat 463 and drought (Barriopedro et al., 2011; Spinoni et al., 2015). As seed bank persistence - often 464 coupled with shorter plant lifespans (Gioria et al., 2020; Thompson et al., 1998) – is theoretically 465 more beneficial for populations in areas with higher environmental variability on the short term 466 (Snyder, 2006), the capacity of a species to produce a persistent seed bank would therefore be a 467 more beneficial strategy in these warmer and more extreme climates with higher community 468 turnover (Childs et al., 2010). Indeed, studies from arid regions have shown seed bank composition 469 470 to be very stable over time, despite natural and experimental climatic variation (DeMalach et al., 2021; Loydi & Collins, 2021). When considering additional functional traits, our analyses 471 supported this suggestion, with annual species more likely to have a higher seed bank persistence, 472 and communities containing a higher relative fraction of annual species also associated with higher 473

474 values of community mean temperature index (Table S3 and Table S5). These analyses also supported previous findings that larger-seeded species are generally found to be more common in 475 warmer regions (Moles & Westoby, 2003; Pakeman et al., 2008), as we showed that higher mean 476 values of seed mass were identified in communities with a higher community mean temperature 477 index. On the other hand, there is evidence that smaller-seeded species generally have higher seed 478 bank persistence (Gioria et al., 2020; Hodkinson et al., 1998), which might be expected to 479 contradict the climatic pattern. However, our species-level analyses found no relationship between 480 seed bank persistence and seed size, while including additional traits in our models did not alter our 481 482 findings that seed banks had a higher community mean temperature index than the herb layer.

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The hypothesis that species that experience a higher level of environmental variability across their 484 485 range should benefit from the ability to form persistent seed banks was also reflected in our finding that climatically widespread species were found to have a higher seed bank longevity (Figure 1b). 486 487 Thus, given that establishment of new individuals from the seed bank is an important filter for community assembly (Larson & Funk, 2016; Marteinsdóttir, 2014), it follows that there will be an 488 accumulation of species with relatively warm and wide climatic ranges in the seed bank. This is 489 reflected in our results that overall, as seed bank communities had a higher community mean 490 temperature index and community temperature range index than the herb layer, even after 491 considering mean values of other species traits that may be related to seed bank persistence. This 492 means that despite common findings of seed bank communities reflecting historical land use and 493 management (Karlík & Poschlod, 2014; Vandvik & Goldberg, 2006), and the potential buffering of 494 climate effects on seed bank richness (Plue et al., 2021), seed bank composition (in our study based 495 on species presence or absence) was not found to lag behind the herb layer in terms of species' 496 climatic associations and are therefore not likely to contribute to slow responses of plant 497 communities to climate warming (Auffret & Thomas, 2019; Bertrand et al., 2011). 498

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500 Our findings are also linked to another aspect of global change and its effects on plant communities. Trends of taxonomic homogenisation at multiple spatial scales despite increases or no net change in 501 species richness (Finderup Nielsen et al., 2019; Keith et al., 2009) imply increases in widespread, 502 503 generalist species that, as well as adding species to communities, are also replacing smaller-ranged specialists over time (Staude et al., 2020, 2022). We found that climatically widespread species are 504 commonly found in the seed bank, and might therefore be playing a role in such community shifts 505 whereby anthropogenic disturbance and climatic warming are actively facilitating their 506 establishment. Indeed, we found that in cooler climates, the effect of widespread species in forming 507 relatively warmer communities in the seed bank compared to the herb layer was smaller (Table 1). 508 That is, the potential for widespread species in the seed bank to contribute to community change has 509 not yet been fulfilled, perhaps due to habitat filtering. Another line of evidence relating widespread, 510 511 seed banking species to recent community change is that seed bank persistence is a trait that has been linked to the successful - and sometimes problematic - establishment of species in new 512 regions (Gioria et al., 2021). Short-lived species are generally associated with ruderal life strategies 513 and often colonise new areas following environmental change (Herben et al., 2018; Pierce et al., 514 2017). Our analyses showed that annual species are more likely to have higher seed bank 515 persistence, and that at the community level are also associated with communities characterised by 516 more widespread species. 517

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This study provides novel insights into the complex role that soil seed banks may play in plant community change in the Anthropocene. In contrast to our initial expectations, seed bank composition did not reflect cooler (potentially past) climatic conditions, but instead contained species adapted to warmer climate than the corresponding herb layer. This was consistent across our climatic gradient and robust to the consideration of additional plant functional traits related to seed bank persistence. We had also hypothesised that habitat would also play a role in the differences in community-level climate associations, related to previous findings of slow community responses to 526 climate change in forests (Zellweger et al., 2020). Following our finding that seed banks contained on average warmer-associated species, an alternative hypothesis could be that forest communities 527 might have larger differences in community mean climate indices between the seed bank and herb 528 529 layer, as cool-related species persist in the herb layer and there is a lack of opportunity for disturbed soil and establishment from the (warmer) seed bank. However, we found instead that grassland 530 habitats exhibited relatively larger differences in community mean temperature indices between the 531 seed bank and herb layer, while forests (which are generally more resistant to invasion than 532 disturbed habitats; Chytrý et al., 2008; Vilà et al., 2007) were found to exhibit a weaker effect of 533 534 widespread species in contributing to warmer seed banks (negative interaction term, Table 1). 535

In sum, rather than contribute to the slow responses of plant communities to climate change, our 536 537 results indicate that warm-associated species with persistent seed banks may play an important role in facilitating community-level responses to climate warming in the future. However, warmer seed 538 bank communities are not likely to reflect a situation where seed banks, but not herb layers, are able 539 to respond to ongoing climate change. Instead, the pattern is probably a reflection of widespread 540 generalists that naturally accumulate in the soil. Therefore, any contribution of the seed bank to 541 community responses to climate change in the herb layer may also contribute to taxonomic 542 homogenisation through the replacement of cooler, range-restricted species. Finally, we also show 543 that by being relatively poor at forming seed banks, cool-associated species may be doubly at risk in 544 a changing climate, both to the long-term shift to a warmer climate and to short periods of extreme 545 conditions that may result in local extinctions. 546

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549 Data availability

550 This study made use of three existing, published databases. Seed bank longevity was calculated 551 from the LEDA traitbase (Kleyer et al., 2008; https://uol.de/en/landeco/research/leda), species

552	climatic indices were taken from ClimPlant (Vangansbeke et al., 2021a, 2021b; now updated to
553	include species calculated for this study), and seed bank and herb layer community data were taken
554	from Plue et al. (2020, 2021).
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565	Author contributions
565 566	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA,
565 566 567	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA, PV, PDF, IA, SB, UG, HJ, AJ, RK, MK, RM, BM and MW digitised range maps for species. AGA,
565 566 567 568	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA, PV, PDF, IA, SB, UG, HJ, AJ, RK, MK, RM, BM and MW digitised range maps for species. AGA, IA, SB, UG, HJ, AJ, RK, MK, RM, BM, MW, RMB, HHB, GD, MH, MJ-B, PM, IEM, RJP, GKP,
565 566 567 568 569	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA, PV, PDF, IA, SB, UG, HJ, AJ, RK, MK, RM, BM and MW digitised range maps for species. AGA, IA, SB, UG, HJ, AJ, RK, MK, RM, BM, MW, RMB, HHB, GD, MH, MJ-B, PM, IEM, RJP, GKP, KT, VV and JP provided data and expertise. AGA led the writing together with JP, PV and PDF.
 565 566 567 568 569 570 	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA, PV, PDF, IA, SB, UG, HJ, AJ, RK, MK, RM, BM and MW digitised range maps for species. AGA, IA, SB, UG, HJ, AJ, RK, MK, RM, BM, MW, RMB, HHB, GD, MH, MJ-B, PM, IEM, RJP, GKP, KT, VV and JP provided data and expertise. AGA led the writing together with JP, PV and PDF. All authors read, commented and approved the manuscript.
 565 566 567 568 569 570 571 	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA, PV, PDF, IA, SB, UG, HJ, AJ, RK, MK, RM, BM and MW digitised range maps for species. AGA, IA, SB, UG, HJ, AJ, RK, MK, RM, BM, MW, RMB, HHB, GD, MH, MJ-B, PM, IEM, RJP, GKP, KT, VV and JP provided data and expertise. AGA led the writing together with JP, PV and PDF. All authors read, commented and approved the manuscript.
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 565 566 567 568 569 570 571 572 573 	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA, PV, PDF, IA, SB, UG, HJ, AJ, RK, MK, RM, BM and MW digitised range maps for species. AGA, IA, SB, UG, HJ, AJ, RK, MK, RM, BM, MW, RMB, HHB, GD, MH, MJ-B, PM, IEM, RJP, GKP, KT, VV and JP provided data and expertise. AGA led the writing together with JP, PV and PDF. All authors read, commented and approved the manuscript.
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 565 566 567 568 569 570 571 572 573 574 575 	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA, PV, PDF, IA, SB, UG, HJ, AJ, RK, MK, RM, BM and MW digitised range maps for species. AGA, IA, SB, UG, HJ, AJ, RK, MK, RM, BM, MW, RMB, HHB, GD, MH, MJ-B, PM, IEM, RJP, GKP, KT, VV and JP provided data and expertise. AGA led the writing together with JP, PV and PDF. All authors read, commented and approved the manuscript. References Alexander, J. M., Chalmandrier, L., Lenoir, J., Burgess, T. I., Essl, F., Haider, S., Kueffer, C., McDougall, K., Milbau, A., Nuñez, M. A., Pauchard, A., Rabitsch, W., Rew, L. J., Sanders,
 565 566 567 568 569 570 571 572 573 574 575 576 	Author contributions AGA, JP, PDF and PV conceived and designed the study. AGA, JP and PV collated data. AGA, PV, PDF, IA, SB, UG, HJ, AJ, RK, MK, RM, BM and MW digitised range maps for species. AGA, IA, SB, UG, HJ, AJ, RK, MK, RM, BM, MW, RMB, HHB, GD, MH, MJ-B, PM, IEM, RJP, GKP, KT, VV and JP provided data and expertise. AGA led the writing together with JP, PV and PDF. All authors read, commented and approved the manuscript. References Alexander, J. M., Chalmandrier, L., Lenoir, J., Burgess, T. I., Essl, F., Haider, S., Kueffer, C., McDougall, K., Milbau, A., Nuñez, M. A., Pauchard, A., Rabitsch, W., Rew, L. J., Sanders, N. J., & Pellissier, L. (2018). Lags in the response of mountain plant communities to climate

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SUPPLEMENTARY INFORMATION

More warm-adapted species in soil seed banks than in herb layer plant communities across Europe

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Figure S1. Map of Europe, showing locations of 54 component datasets providing seed bank and herb layer plots for the seed bank dataset used in this study. All plots within each dataset are categorised along a land-cover gradient from open grassland habitats (blue squares) over transitional habitats with developing canopies (yellow circles) towards mature, closed canopy forest habitats (green triangles). Figure reproduced from Plue et al. (2021; CC BY 4.0 License).



Figure S2. Scatterplot of the species mean temperature index (SMTI; mean temperature experienced by a species across its European range) and the species temperature range index (STRI; the difference between the 5th and the 95th percentile of temperatures experienced by a species across its European range), for 840 species for which species temperature indices exist and seed bank longevity index could be calculated. The red (upper-left) point shows *Sedum anglicum*, which has an SMTI of 11.0°C and an STRI of 4.0°C. The blue (lower-left) point shows *Stellaria borealis*, which has a mean of SMTI of 0.3°C and STRI of 4.9°C. These examples illustrate that species with narrow climatic ranges in Europe are not necessarily cool-associated, as also illustrated by a low correlation (-0.1) between SMTI and STRI.

Table S1. Main characteristics of the component studies of the seed bank database. Plots per category refers to the number of plots								
from each dataset within each of the following categories: 1 = Open, grassland habitats, 2 = Abandoned grassland and recent forest								
habitats, characterized by a developing forest canopy, 3 = Mature, old-growth forest habitats with fully developed, closed canopy,								
		Plots per category Herb layer		Herb layer	Seed bank			
Dataset	Region	1	2	3	Plot area [m ²]	Collection	Plot area	Plot depth
Auffrant & Coursing (2011), https://doi.org/10.1111/j.1265.2664.2011.02010.v)	C Sweden	400	FFF		1	year	[m²]	[m]
Autret & Cousins (2011; <u>mtps://doi.org/10.1111/j.1385-2664.2011.02019.x</u>)	C Sweden	433	555		0.05	2006	0.0064	0.07
Adestad et al. (2015; <u>https://dx.doi.org/10.1007/S11256-013-0207-x</u>)		100			0.25	2005	0.003927	0.05
Basio et al. (2015, <u>https://doi.org/10.1036/fic0111157165</u>)	England	30			4.5	2009	0.000362	0.15
Berker et al. (1999; <u>https://doi.org/10.1046/j.1365-2745.2000.00485.x</u>)	Netherlands	80	00	4.4	25	1992	0.01257	0.1
Bossuyt et al. (2002; https://doi.org/10.1023/A:1020391430072)	Beigium		23	11	1	1999	0.01924	0.2
Calcada et al. (2015; <u>https://doi.org/10.1007/s10980-015-0236-y</u>)	N France			92	4	2008	0.00628	0.1
Decocq et al. (2004; <u>https://doi.org/10.1023/B:BIOC.0000048454.08438.c6</u>)	N France			10	400	1999	0.02512	0.2
Decocq & Calcada (unpublished)	N France			10	4	2008	0.00628	0.1
Devlaeminck et al. (2005; <u>https://doi.org/10.1007/s11258-004-0008-2</u>)	Belgium			35	1	2002	0.05772	0.2
Donelan and Thompson (1980; <u>https://doi.org/10.1016/0006-3207(80)90029-4</u>)	England		3	1	300	1978	0.2308	0.07
Dutoit & Alard (1995; <u>https://doi.org/10.1007/BF00058205</u>)	N France	2	2		2	1993	0.07691	0.085
Grandin & Rydin (1998; <u>https://doi.org/10.1046/j.1365-2745.1998.00252.x</u>)	C Sweden		63		1	1993	0.00152	0.03-0.15
Jacquemyn et al. (2011; <u>https://doi.org/10.1016/j.biocon.2010.09.020</u>)	Belgium	10	5		1	2009	0.009813	0.1
Jakobsson et al. (2006; <u>https://doi.org/10.1139/b06-136</u>)	C Sweden	100			1	2001	0.005	0.07
Jankowska-Blaszczuk et al. (1998; https://doi.org/10.1023/A:1009750201803)	Poland			50	4	1995	0.0961	0.05
Kalamees & Zobel (1997; <u>https://doi.org/10.1007/BF02803879</u>)	Estonia	6	6		225	1993	0.0577	0.05
Kalamees et al. (2012; <u>https://doi.org/10.1111/j.1654-109X.2011.01169.x</u>)	Estonia	4	4		100	2005	0.1	0.05
Kalamees & Zobel (2002; <u>https://doi.org/10.1890/0012-</u> 9658(2002)083[1017:TROTSB]2.0.CO:2)	Estonia	10			10	1996	0.1556	0.1
Koch et al. (2011; <u>https://doi.org/10.1016/j.flora.2010.01.010</u>)	N Germany	3			16	2005	0.03533	0.2
Lee et al. (2013; <u>https://doi.org/10.1016/j.biocon.2013.05.023</u>)	England	24			1	2010	0.009817	0.07
Marage et al. (2006; <u>https://doi.org/10.1139/b05-142</u>)	S France		8	8	400	2001	0.01768	0.05
Marrs (unpublished)	England	50			5	2003	0.005655	0.07
Marrs (unpublished)	England	47			5	2003	0.005655	0.07
Marteinsdottir (2014; https://doi.org/10.1371/journal.pone.0103352)	C Sweden	12			100	2011	0.02513	0.05
Milberg & Hanson (1994; <u>https://doi.org/10.2307/3235635</u>)	S Sweden	10	10		1	1990	0.01282	0.1
Måren & Vandvik (2007; <u>https://doi.org/10.1111/j.1654-1103.2009.01091.x</u>)	Norway	40			1	2004	0.009815	0.06
Pakeman (unpublished)	Scotland	66			4	2006	0.01767	0.1
Pakeman & Hay (<u>https://doi.org/10.1006/jema.1996.0057</u>)	England	18			1	1993	0.008482	0.1
Pakeman & Marshall (<u>https://doi.org/10.1046/i.1365-2699.1997.00127.x</u>)	England	38			12.5	1994	0.175	0.05
Pakeman et al. (https://doi.org/10.1046/j.1365-2745.1998.00327.x)	England	32			0.25	1996	0.01	0.05
Plue et al. (2008; https://doi.org/10.1007/s10980-008-9229-4)	N France	1		46	100	2004	0.01328	0.2

Plue et al. (2009; https://www.jstor.org/stable/pdf/20794668.pdf)	Belgium			35	100	2006	0.01125	0.2
Plue et al. (2010; <u>https://doi.org/10.1017/S0960258509990201</u>)	N France			5	4.41	2006	0.1232	0.05
Plue et al. (2010; <u>https://doi.org/10.1016/j.ppees.2010.03.001</u>)	N Germany			119	100	2007	0.01306	0.1
Plue & Hermy (2012; <u>https://doi.org/10.1111/j.1654-1103.2011.01361.x</u>)	Belgium	3			4.41	2001	0.1232	0.05
Plue & Cousins (2013; <u>https://doi.org/10.1016/j.biocon.2013.02.010</u>)	C Sweden	48	86		4	2010	0.02405	0.05
Plue et al. (2013; <u>https://doi.org/10.1111/geb.12068</u>)	France - Sweden		26	118	4	2011	0.0481	0.1
Plue & Cousins (2018; <u>https://doi.org/10.1111/oik.04813</u>)	C Sweden	77			4	2014	0.0481	0.05
Thompson (1986; https://doi.org/10.2307/2260394)	England	1			0.6272	1982	0.6272	0.05
Thompson et al. (1994; <u>https://doi.org/10.2307/2390108</u>)	England	1			0.6272	1985	0.6272	0.05
Van Calster et al. (2008; <u>https://doi.org/10.3170/2008-7-18405</u>)	M France			48	400	2005	0.009891	0.1
Vanderhasselt & Plue (unpublished)	Belgium		93		3	2008	0.01924	0.1
Vandvik & Goldberg (2006; <u>https://doi.org/10.1086/505759</u>)	Norway	12	6		4	1997	0.03768	0.1
Vandvik et al. (2015; <u>https://doi.org/10.1111/oik.02022</u>)	Norway	26			0.41	2008	0.41	0.03
Wagner et al. (2003; https://www.jstor.org/stable/23726882)	Estonia	2	2		100	1998	0.1256	0.12
Warr et al. (1994; <u>https://doi.org/10.2307/2845469</u>)	England			20	180	1990	0.0491	0.15

Table S2. Pearson correlation coefficients between the different traits used to predict seed bank longevity in the auxiliary analyses. For life span, 0=annual or biennial, 1=perennial. SMTI=Species mean temperature index; STRI=Species temperature range index.

	Life span (binary)	Seed number	Seed mass (mg)	SMTI (°C)	STRI (°C)
Life span (binary) Seed number Seed mass (mg) SMTI (°C) STRI (°C)	0.05 0.04 -0.15 -0.09	0.05 -0.003 0.03 0.06	0.04 -0.003 0.14 -0.06	-0.15 0.03 0.14 -0.1	-0.09 0.06 -0.06 -0.1

Table S3. Species-level predictors of seed bank longevity index using binomial linear models. Number of species represents the number of species for which each combination of trait data were available for analysis. For life span, 0=annual or biennial, 1=perennial. SMTI=Species mean temperature index; STRI=Species temperature range index.

	Estimate	Std. Error	t-value	P.value			
Species mean temperature index (840 species; Model 1a)							
Intercept	-1.66	0.21	-7.85	<0.001			
SMTI (°C)	0.12	0.03	4.66	<0.001			
Species temperature range index (840 species; Model 1b)							
Intercept	-2.19	0.2	-10.79	<0.001			
STRI (°C)	0.12	0.02	7.6	<0.001			
SMTI and all other traits (511 species; Model	1c)						
Intercept	-0.81	70260	0	1			
SMTI (°C)	0.14	7734	0	1			
Seed mass (mg)	-0.84	6649	0	1			
Seed number	<0.001	0.07	0	1			
Lifespan (binary)	-0.77	29390	0	1			
STRI and all other traits (511 species; Model	1d)						
Intercept	-1.0	60220	0	1			
STRI (°C)	0.1	4165	0	1			
Seed mass (mg)	-0.17	6457	0	1			
Seed number	<0.001	0.002	0	1			
Lifespan (binary)	-0.91	28940	0	1			
Seed mass (804 species; Model 1e)							
Intercept	-0.42	9540.64	0	1			
Seed mass (mg)	-0.17	4594.57	0	1			
Seed number (580 species; Model 1f)							
Intercept	-0.06	0.005	-12.88	<0.001			
Seed number	<0.001	<0.001	-0.032	0.98			
Lifespan (717 species; Model 1g)							
Intercept	-0.038	0.11	-0.36	0.72			
Lifespan (binary)	-0.9	0.12	-7.6	<0.001			

Table S4. Pearson correlation coefficients between the community mean values of different traits used as control variables for the auxiliary analysis of differences in community climate indices between seed bank and herb layer

	Life span (binary)	Seed number	Seed mass (mg)
Life span (binary)		0.022	0.12
Seed number	0.022		-0.14
Seed mass (mg)	0.12	-0.14	

Table S5. Linear mixed effect model outputs for community-level analysis of climate indices in relation to source (seed bank or herb layer), MAT (mean annual temperature in °C) and functional traits. For life span, 0=annual or biennial, 1=perennial.

	Estimate	Standard	t-value	95% CI			
Oliveration and the Lafe and the		Error					
Climatic control of community mean temperature indices, plus differences across sources							
(Model 2a)	5.04	0.40	00.40	E 4 E 10 4 0			
Intercept	5.84	0.18	32.48	5.45 - 6.16			
MAT (°C)	0.2	0.02	9.69	0.16 – 0.24			
Source (herb layer)	-0.23	0.06	-4.05	-0.36 – -0.13			
MAT:Source (herb layer)	0.003	0.007	-0.39	-0.01 – 0.01			
Differences across sources for community temperature range indices (Model 2b)							
Intercept	12.43	0.12	107.79	12.21 - 12.66			
Source (herb layer)	-0.64	0.02	-25.65	-0.68 – -0.59			
Differences across sources for community mean temperature indices, controlling for mean							
community values of funct	ional traits (Mod	lel 2c)					
Intercept	8.05	0.13	71.33	7.8 – 8.28			
Source (herb layer)	-0.18	0.016	-11.23	-0.21 – -0.15			
Seed mass (mg)	0.026	0.003	9.31	0.02 – 0.031			
Seed number	<0.001	<0.001	15.74	<0.001 - <0.001			
Lifespan (binary)	-0.78	0.06	-12.94	-0.90.66			
Differences across sources for community temperature range indices, controlling for mean							
community values of functional traits (Model 2d)							
Intercept	12.81	0.15	86.91	12.49 – 13.08			
Source (herb layer)	-0.53	0.029	-18.49	-0.59 – -4.74			
Seed mass (mg)	-0.068	0.005	-13.24	-0.078 – -0.058			
Seed number	<-0.001	<-0.001	-6.39	<-0.001 - <-0.001			
Lifespan (binary)	-0.27	-0.14	-2.4	-0.480.06			