Dissecting the hydrological niche: soil moisture, space and lifespan

Running title: Dissecting the hydrological niche

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Summary

1. Niche segregation on fine-scale soil-moisture gradients is found in many plant communities, but the contribution of this to community structure has yet to be analysed in a fully spatial manner. We introduce a univariate and multivariate analytical approach that, taking spatial autocorrelation into account, decomposes the spatial structure of species composition into components that correlate with linear and non-linear soil-moisture gradients plus a residual.

2. The analysis is applied to two contrasting plant communities in Western and Central-Northern Spain, a seasonally dry dehesa meadow that is rich in both annual and perennial herbs and an alpine meadow community. In each we: (i) determined whether the community was structured on a soil moisture gradient, (ii) tested whether there was spatial segregation into hydrological niches, (iii) characterised the hydrological niches of individual species, and (iv) controlling for spatial autocorrelation, determined how much of the spatial structure in the community was due to variation in hydrology. In the dehesa community we also compared annuals and perennials with respect to questions 1-4.

3. We found that both plant communities were primarily structured along hydrological gradients and that spatial segregation into hydrological niches occurred among perennial species, though not among the annuals in the dehesa community. Dehesa annuals were spatially aggregated in the driest niches. Hydrological variation shaped the responses of 60% of the annual and about 70% of the perennial species in both the dehesa meadow and the alpine community. Most responses were either monotonic or hump-shaped. Finally, spatially structured hydrological variation proved to be the main
driver of spatially structured species composition in all three cases.

4. Synthesis Consistent univariate and multivariate results showed that linearly (gradient of slope) and topographically (at a fine scale) structured variation in hydrology is the main driver of spatially structured species composition in both test communities. Our results support the hypothesis that spatial niche segregation on soil-moisture gradients is an important mechanism of coexistence for perennials in both test communities, though not for the species-rich sub-community of annuals in the dehesa meadow.

Key words: dehesa, RDA models, hydrological niche, Iberian Peninsula, GAMM regression, Pianka’s index, plant coexistence, Water Table Depth, MEM spatial variables
Introduction

Most plant communities contain mixtures of species that compete for the same essential resources. How competing plants manage to coexist with one another is a long-standing conundrum (Silvertown 2004), but we may at last be nearing a solution. The various mechanisms of coexistence that have been proposed can be divided into two types (Chesson 2000): stabilizing mechanisms such as niche segregation, in which the effects of interspecific competition are frequency-dependent, thus protecting species from local extinction when they become rare, and equalizing mechanisms such as the neutral theory (Hubbell 2001), that limit or delay the monopolization of resources by potentially dominant species.

For a decade after the publication of Hubbell's (2001) book, *The unified neutral theory of biodiversity and biogeography*, it was often argued that plant communities must be assembled by equalizing mechanisms because the plant niches that would stabilize communities had not been demonstrated (Rosindell, Hubbell & Etienne 2011). Silvertown (2004) pointed out that an absence of evidence for niche-based coexistence was not evidence of its absence. Purves & Turnbull (2010), showed that the central assumption of neutral theory, which is that species that are different in phenotype will have equal fitness, is only likely to be true in the rarest of circumstances.

Examples of stable coexistence achieved through niche segregation and tested in competition models have now begun to accumulate (Adler et al. 2006; Angert et al. 2009; Levine & HilleRisLambers 2009; Adler, Ellner & Levine 2010; Clark et al. 2010). Nonetheless, the ultimate solution to the conundrum of coexistence is likely to be pluralistic because it is widely recognised that the composition of plant communities can be influenced by both stabilizing and equalizing mechanisms to varying degrees (Adler, HilleRisLambers & Levine 2007; Stokes &
Archer 2010; Chase & Myers 2011; Rosindell et al. 2012; Chase 2014), although stabilizing mechanisms of some kind are essential for indefinite coexistence (Chesson 2000). Wilson (2011) evaluated the 12 theories that he believed contain the only distinct mechanisms of plant coexistence and concluded that 5 stabilizing mechanisms and 2 equalizing ones had at least some empirical support. Niche segregation is the best-supported stabilizing mechanism and there is growing field and experimental evidence that soil-moisture gradients are an important niche dimension in many plant communities (Silvertown et al. 1999; Araya et al. 2011; Markham 2014), reviewed by Silvertown, Araya & Gowing (2014).

While segregation on soil-moisture gradients appears to be ubiquitous across the gamut of plant communities from arid environments through to wetlands (Silvertown, Araya & Gowing 2014), we still do not know what contribution this makes to plant community structure or coexistence. In this paper we introduce a new methodology that makes it possible to answer the first of these questions. We use this methodology to dissect the hydrological niche in two different plant communities and to estimate how much of the spatial variance in plant community structure is due to segregation on a soil-moisture gradient and how much is due to other processes including spatial autocorrelation. Both plant communities are in Spain, one in a wet, sub-alpine environment containing only perennial herbs and the other is a lowland, seasonally dry dehesa grassland with a high diversity of both annuals and perennials.

Annuals and perennials have different regeneration biology, with possible consequences for coexistence (Grubb 1977). Many annuals have life cycles that contain a persistent seed bank, which lends itself to coexistence mediated by temporal niche segregation and the storage effect (Warner & Chesson 1985; Pake & Venable 1996; Angert et al. 2009). This might mean that spatial niche segregation is weaker in annuals than in perennials and so we also test for this.
Our analysis enables us to answer five questions:

1. Are the communities structured on a hydrological (soil moisture) gradient?
2. Is there spatial segregation into hydrological niches?
3. What is the shape of the hydrological niches of individual species?
4. Controlling for spatial autocorrelation, how much of the spatial structure in the community is due to variation in hydrology?
5. Do annuals and perennials behave alike with respect to questions 1 - 4?

Materials and methods

Sampling design

An Iberian dehesa meadow (Eunis habitat type 6310) (European Commission 2007) on granite soils was sampled at La Mina in Moscosa Farm (41º 8' 21.88'' N, 6º 6' 52.33'' W; 780 m a.s.l.), Salamanca province, Western Spain. A 50 x 50 m study plot with a 1.8% gradient was sampled (Appendix S1). The presence of plant species was recorded (Spring 2007) in 196 1-m² quadrats placed on a 14 x 14 grid. Similarly, an Iberian alpine meadow on sandstone/conglomerate soils, with a 2.1% gradient, was sampled at Laguna Larga in the Urbión Peaks (42º 0' 19.50'' N, 2º 52' 2.26'' W; 2080 m a.s.l.), Soria province, Central-Northern Spain, using 172 1-m² quadrats placed regularly. Plant nomenclature followed standard Floras (Tutin et al. 1964-1980; Castroviejo 1986-2011), except for the species included in Appendix S1. The spatial variables northing, easting and elevation were measured using a total station machine (Leica Geosystems TPS800).
Quantification of the hydrological gradient

We made fortnightly measurements of water-table depth from nine dipwells over a two-year period. A fine-scale topographic map (constructed from the surveyed points) was used to construct a field-scale hydrological models for each of the sites (Gowing & Youngs 1997). This model quantified by interpolation the average water-table depth (AWTD) in each quadrat during the growing season (30 weeks for Moscosa, from mid February to end of September; 20 weeks for Urbión, from mid May to mid September). Since water-table depth measurements are made from an origin at ground level, low values of AWTD correspond to high levels of oxygen-deficit stress (due to waterlogging); high values of AWTD correspond to high levels of water-deficit stress (due to soil drying) over the growing season.

Data analysis

Our dissection of the spatial distribution of species in relation to soil-moisture gradients had four steps. First, we used unconstrained non–metric multidimensional scaling (NMDS) to reduce the dimensionality of the plant distribution data without any reference to environmental gradients of any kind. We then tested whether the principal dimensions that result from the NMDS analysis align with the soil-moisture gradient. In the second step, we tested for niche segregation against a null model; in the third step we characterised the hydrological niche of each species using Generalised Additive Mixed Models (GAMMs) while taking spatial autocorrelation into account. Finally, in the fourth step, we partitioned the spatial variance in species distribution into three components, (i) an hydrological component, (ii), a linear trend component, and (iii) and a spatial component defined through sets of independent spatial variables constructed using the Moran’s eigenvector maps method.
At both sites, we used a two-matrix dataset. For Moscosa, it included a species composition matrix of $n = 196$ sample units x $p = 123$ species, where each element represented the presence-absence of a species in a sample unit. An environment and spatial matrix of $n = 196$ sample units x $q = 3$ represented the values of AWTD, northing, and easting. The same matrices were used for Urbión, with $n = 172$ sample units and $p = 52$ species.

**Step 1: reducing dimensionality**

We explored the main underlying gradients in the species composition of the communities by means of unconstrained ordination, i.e. by species ordination determined only by the species data and not by any other external (environmental) variables. For this purpose, non–metric multidimensional scaling (NMDS) (Kruskal 1964; McCune & Grace 2002) was used because this technique preserves the ordering among objects and may be used on the basis of any distance matrix (Legendre & Legendre 2012). NMDS was here performed on the basis of the Jaccard dissimilarity measure (Legendre & Legendre 2012), which is directly interpretable as a (dis)similarity percentage and does not take joint absences into account (Field, Clarke & Warwick 1982). Prior to NMDS itself, in order to reduce the data dimensionality (McCune & Grace 2002) and hence facilitate a simplified ordination in two dimensions, 42 species with relative frequency $\leq 0.05$ were deleted from the Moscosa species matrix and 24 from the Urbión species matrix. Then the ordination was performed through the function metaMDS() of the package vegan (Oksanen *et al.* 2013) in R software (R Core Team 2013). The correct preservation of the ordering relationships among objects was assessed by means of a Shepard diagram (Borcard, Gillet & Legendre 2011); the non-metric goodness-of-fit of the ordination were $R^2 = 0.967$ (Moscosa) and $R^2 = 0.989$ (Urbión). We then used the function ordisurf() in vegan to fit a quadratic trend surface corresponding to the hydrological descriptor AWTD onto
the ordination that had been defined. As a result, the AWTD surface thus overlaid onto the NMDS ordination did not influence the ordination itself.

**Step 2: testing for niche segregation**

We computed pairwise values of Pianka’s index of niche overlap (Pianka 1973) for all species combinations to test the null hypothesis of random overlap between species. Indices were computed by dividing the available niche space into sections of AWTD width equal to 0.05 m. Hence we defined thirteen hydrological niches: AWTD (m) = 0.200-0.249 (first niche), 0.250-0.299 (second niche), …, 0.800-0.849 (thirteenth niche) for the Moscosa site, but only seven niches (0.200-0.249, 0.250-0.299, …, 0.500-0.549 ) for the Urbión site because its hydrological gradient is shorter. AWTD was measured in a positive downwards direction with origin at ground level. Hence the severity in conditions of oxygen-deficit stress in the first niche is at a maximum (due to soil flooding), whereas the severity in conditions of water-deficit stress is at a minimum. Conversely, the severity in conditions of water-deficit stress in the thirteenth niche is at a maximum (due to soil drying), whereas the severity in conditions of oxygen-deficit stress is at a minimum. The availability of the different resource states was introduced into the analysis (Hurlbert 1978). Departures of mean niche overlap from random expectation were determined by a randomization test that used 10,000 simulated matrices (“RA4 randomization algorithms”) in EcoSim 7.0 software (Gotelli & Entsminger 2009); the p-values reported in the results section correspond to the probability that the simulated index ≤ the observed index.

**Step 3: characterising hydrological niches**

Generalised Additive Mixed Modelling (GAMM regression) was used to model the relationships between the presence of individual species and the hydrological descriptor AWTD, while controlling for spatial autocorrelation to improve estimation and inference.
Tests were restricted to 81 (Moscosa) and 28 (Urbión) species with relative frequency > 0.05, for which the null hypothesis that AWTD has no effect on the probability of presence for species was tested. This non-parametric model is, in summary, the additive modelling equivalent of a logistic regression model (Zuur et al. 2010). Additive modelling fits smoothers (smoothing curves) through the data without assuming linearity and is thus able to describe a wide variety of relationships (Wood 2006). Additive modelling was implemented here via the function gamm() of the package mgcv (Wood 2011). This function was chosen because it automatically determines the right amount of smoothing and it allows mixed models to be used for including spatial correlation structure (Pinheiro & Bates 2000), as may be required when autocorrelation is present in model residuals.

Additive modelling assumes independence, but there are at least two reasons for expecting spatial autocorrelation (SAC) (Griffith 1992) in our species data. First, SAC will occur if the habitat is spatially structured, i.e. if the habitat variables that drive species are themselves spatially correlated. Second, SAC may occur because of spatially constrained species dispersal. In the first case, we expect environmental explanatory variables to partially account for this SAC; however, since it is unlikely that the presence of species is driven by hydrology alone, we may also expect some remaining SAC in model residuals. SAC due to species dispersal will not be accounted for by environmental variables. In summary, the presence of SAC must be assessed for each species model by means, for example, of spline correlograms (Bjornstad & Falck 2001) that are here estimated using the function spline.correlog() of the package ncf (Bjornstad 2013). Therefore, for each species, if no SAC was found in the residuals of the basic model (as defined above), we proceeded to validate the fitted model as usual (Zuur, Ieno & Smith 2007) and then employ it for statistical inference. If, on the contrary, SAC was found in the model residuals (i.e. if the assumption of independence did not hold), we identified an
appropriate spatial autocorrelation structure (Pinheiro & Bates 2000) to account for SAC and then re-fitted the model. Appropriate structures were identified using semivariograms (Cressie 1993), which was implemented here with the function Variogram() of the package nlme (Pinheiro et al. 2014). Once identified, we introduced the appropriate spatial autocorrelation structure into the basic model (Zuur et al. 2010) and re-fitted it using the above function gamm(). Though, as stressed above, the effect of AWTD on the presence-absence of species was fitted while controlling for SAC, the results presented in Tables 1-3 and Figures 2-4 focus on the models’ fixed parts (the smoothers) because this work deals with such (hydrological) effects.

**Step 4: partitioning spatial variance**

Redundancy analysis, or RDA (Wollenberg 1977; Legendre & Legendre 2012), was used to test for relationships between species composition (response) and spatial or hydrological predictors. RDA can be described as a technique for multivariate regression (Zuur, Ieno & Smith 2007). The response was a Hellinger-transformed (Legendre & Gallagher 2001) presence-absence species matrix. We used three kinds of predictors: (i) MEM spatial variables (see below); (ii) a linear trend; and (iii) the hydrological predictor AWTD. The linear trend is a surface specified by the X-Y coordinates and is here used to describe linear spatial structures. In order to model non-linear hydrological relationships (Borcard, Gillet & Legendre 2011), the hydrological component included not only an AWTD first-degree term, but also second- and third-degree terms; these were computed using the function poly() of the package stats in R software (R Core Team 2013), so they are orthogonal. The spatial component was modelled using sets of independent spatial variables that were specifically constructed for each species assemblage using the MEM (Moran’s eigenvector maps) method (Dray, Legendre & Peres-Neto 2006; Legendre & Legendre 2012). These MEM spatial variables, which can model
positive and negative spatial autocorrelation, were constructed using weighted connectivity matrices (Borcard, Gillet & Legendre 2011) via the R package spacemakeR (Dray 2013).

In addition to RDA marginal tests, we carried out a partitioning of variation in species composition (Borcard, Legendre & Drapeau 1992; Legendre & Legendre 2012) into components of unexplained variation, non-spatial hydrological variation and non-hydrological spatial variation (either as a linear surface or as modelled by MEM variables); besides, fractions corresponding to spatial structuring shared by hydrological conditions (induced spatial variation), and others, were also estimated. This partitioning supports models of causal relationship (Legendre & Legendre 2012) and here was used mainly in order to estimate the extent of induced spatial variation, i.e. spatially structured species composition that is explained by spatially structured hydrological variation. Induced spatial variation has two fractions. The first (fraction \([g]\) in the partition diagram) corresponds to spatially structured variation in species composition that is associated with linear gradients, i.e. with the sites’ gradients of slope. The second (fraction \([f]\) in the partition diagram) corresponds to spatially structured variation in species composition that is associated with the sites’ local topographies. The unique contribution of each component, as well as the extent of overlap among them, was estimated by varying the order of fit using the functions rda() and varpart() of the R package vegan (Oksanen et al. 2013).

Supporting information provides data (Appendices S4 and S5) and R coding (Appendix S3) sufficient to replicate the analysis described above.

**Results**

At Moscosa, we recorded 71 species of annual and 52 perennials (Appendix S1), but only 81 of the 123 species had a relative frequency greater than 0.05; in Urbión, 52 species were found,
but only 28 had frequency greater than 0.05 (Appendix S1). Unconstrained ordination (Fig. 1), which here shows the two main species gradients in the meadows without external reference to any environmental variables, shows that the first main species gradient (NMDS1) is, in both cases, strongly associated with the hydrological gradient (as measured by AWTD in m).

For Moscosa, the null hypothesis of random overlap across the hydrological space at a fine scale (thirteen niches) was rejected for the whole community (observed mean = 0.399 < simulated index = 0.410; \( p = 0.000 \)) and for perennials as a group (observed mean = 0.369 < simulated index = 0.383; \( p = 0.001 \)). Hence perennials segregate along the hydrological gradient. Significant GAMM models were fitted for 23 (70% of species with frequency > 5%) perennials (Table S1 in Appendix S2). The average model fit was \( R^2 \) (adj.) = 21.3\%. Consistent with the observed segregation pattern, monotonic perennial responses were either increasing or decreasing with increasing soil drying (Fig. 2), thus contributing to segregation (compare, for example, \( Poa bulbosa \) with \( Poa trivialis \) or \( Senecio jacobaea \) with \( Thapsia villosa \) in Fig. 2).

The various hump-shaped, or similar, responses (35% of the fitted models) also contribute to segregation (compare, for example, \( Briza media \), \( Galium verum \) and \( Echium plantagineum \) in Fig. 2). In contrast and as suggested by the unconstrained ordination (Fig. 1), the null hypothesis of random overlap was not rejected for annuals (observed mean = 0.463 > simulated index = 0.448; \( p = 0.995 \)). Hence annuals do not segregate along the hydrological gradient, but rather tend to aggregate at the dry end. Significant GAMM models were fitted for 26 annuals (54\%) (Table S2 in Appendix S2), with average model fit \( R^2 \) (adj.) = 18.6\%. No significant relationships were found for invasive annuals (e.g. \( Trifolium dubium \)) with relative frequency greater than c.0.9. Consistent with the observed aggregation pattern, most annual responses (60\% of the fitted models) are both monotonic (either sigmoid or curvilinear) and increasing with increasing soil drying (Table S4; Fig. 3). This contributes greatly to generate this pattern.
of species aggregation (compare *Aphanes arvensis*, *Bellardia trixago*, *Brassica barrelieri*, *Galium parisiense*, *Jasione montana*, *Ornithopus perpusillus*, *Trifolium glomeratum* and *Xolantha guttata* in Fig. 3), in spite of 24% of fitted models for annuals being found to display hump-shaped relationships.

For Urbión, the null hypothesis of random overlap across the hydrological space at a fine scale (seven niches) was rejected for the whole community (observed mean = 0.579 < simulated index = 0.595; \( p = 0.010 \)). Hence species segregate along the hydrological gradient. Significant GAMM models were fitted for 20 (71% of species with frequency > 5%) species (Table S3 in Appendix S2). The average model fit was \( R^2 \) (adj.) = 17.2%. Consistent with the observed segregation pattern, monotonic species responses were either increasing or decreasing with increasing soil drying (Fig. 4), thus contributing to segregation. Hump-shaped responses (25% of the fitted models) also contribute to segregation.

Marginal tests (Table 1) show that, for Moscosa, AWTD explains c.18% of multivariate variation in perennials composition and c.16% in annuals composition. Likewise, species composition is strongly spatially structured (c.30% for perennials; c.27% for annuals, as described by MEM spatial variables), with a linear gradient (Table 1; Fig. 5) being responsible for a relatively important part of these spatial structures. Variation partitioning (Fig. 5), however, shows that the unique contribution of the hydrological descriptors to explain composition (fraction [a]) is less than 2% in all three cases, i.e. species composition explained by non-spatially structured hydrological variation is minor. In other words, induced spatial variation (fractions [f] and [g]), which corresponds to spatially structured species composition that is explained by spatially structured hydrological variation, is the strongest element in all three cases. The sum of fractions [f] and [g] amounts to 16.2% of the variance explained in the perennials assemblage and 14.3% in the annuals assemblage. In both cases fraction [g],
corresponding to variation in species composition that is associated with the (linear) gradient of slope, is the most important (9.6%). Fraction \( f \) corresponds to spatially structured variation in species composition that is not associated with linear gradients, but with local topography; this fraction is stronger for perennials (6.6%) than for annuals (4.7%).

For the Urbión meadow, AWTD explains c.16% of multivariate variation in species composition (Table 1). Likewise, species composition is strongly spatially structured (38.7%), as described by MEM spatial variables). Variance partitioning (Fig. 5) shows that species composition explained by non-spatially structured hydrological variation (fraction \( a \)) is minor (2.4%). Hence, as in the Moscosa site, induced spatial variation (fractions \( f \) and \( g \)), which together add up to 11.9%, represents a strong component). However, in the Urbión site, the spatially structured variation in species composition that is associated with the local topography (fraction \( f \)) explains 8.1% of adjusted variance and is, therefore, greater than the variation in species composition that is associated with the (linear) gradient of slope (fraction \( g \)), which represents only 3.8% of adjusted variance.

Finally, in both meadows, the unique contributions of the MEM spatial variables (fraction \( c \)), which correspond to spatially structured species composition that is not explained by the hydrological descriptor, but by latent processes, suggests the existence of spatially structured ecological factor(s) other than AWTD driving species composition. This component is stronger in the Urbión site (14.8% of total variance) than for the annuals (8.3%) or the perennials (10.1%) in the Moscosa site.

**Discussion**

Although soil moisture and local topography are well known influences on plant distribution (Moeslund *et al*. 2013), we believe that this is the first study to formally decompose plant
distribution into spatial components that include the important effect of hydrology. The methods developed by Borcard (1992) and Borcard & Legendre (1994) have been widely used, for example to test competing theories regarding dispersal limitation, environmental determinism and neutral models in an American temperate forest (Gilbert & Lechowicz 2004). In Amazonian forests, Tuomisto et al (2003) found that spatially-structured environmental variation was the most important ecological factor explaining plant composition at a regional scale, with dispersal having also some ecological effect though neutrality was not supported.

By analysing the spatial variance in plant community structure, we have been able to answer 5 questions about the hydrological niche, including whether annuals and perennials behave alike. First, we found that both the plant communities that we investigated were structured along hydrological gradients and that their primary axes of variation aligned with Average Water Table Depth (Fig. 1a, b). Average Water Table Depth (AWTD) is measured as a distance below the surface so this measure of hydrological conditions is necessarily highly correlated with fine scale topography. Second, we found that spatial segregation occurred among perennial species, though not among the annuals in the dehesa community (Fig.1a). Dehesa annuals as a group were aggregated at the dry end of the hydrological gradient where most fell into just three or four niches (niches 0.50-0.70) out of the 17 that were present.

Third, we investigated the shape of species' hydrological niches by fitting GAMS, which showed that species responses were, with very few exceptions, either monotonic (increasing or decreasing along the hydrological gradient) or hump-shaped (between 24-35% of species responses). The important conclusion here is that, contrary to the assumptions of the neutral model, co-occurring species show different responses along soil-moisture gradients. Similar results have now been found in many plant communities (Silvertown, Araya & Gowing 2014).
A comparison of the shapes of the species' hydrological niches illuminates the difference in community structure found between perennials and annuals. Since most perennial responses were either monotonic increasing or monotonic decreasing with increasing soil drying (65-75%), this difference created segregation, with hump-shaped responses also contributing to segregation. In contrast, most annuals responses were monotonic increasing with increasing soil drying (62%), thus creating a pattern of overlap and species aggregation. These results suggest that, at least in the *dehesa* community we studied, spatial niche segregation on soil moisture gradients may not be an important mechanism of coexistence in the species-rich sub-community of annuals. Given that rainfall in the area is highly variable from year-to-year (Ceballos, Morán & López 2013), temporal niche segregation (the storage effect), as found among Sonoran desert annuals by Angert *et al.* (2009), is an alternative possibility.

Fourth, we partitioned the components of spatial structure in the two plant communities, with complex results (Table 1 and Fig. 5). Among the perennials at Moscosa, 18% of the variance in species composition was ascribed to variation in hydrology. For the annuals in Moscosa and for perennials in Urbión, the variance explained was about 16% in both (16.2% and 15.6%, respectively). Nearly all the variance in hydrology that drove species composition was spatially structured, but about 2% of the adjusted variance explaining change in species composition was not. This may simply reflect error in the hydrological models fitted. This spatial structure can be further subdivided into components that correspond to the linear gradients of slope (fraction \([g]\)) and to local topographic variation (fraction \([f]\)). In Moscosa, the gradient of slope (9.6% for both annuals and perennials) was more important than local topography (6.6% for perennials and 4.7% for annuals). In contrast, local topography (8.1%) was more important in Urbión than the gradient of slope (3.8%).
Overall, the contributions of hydrologically-correlated spatial variation may appear rather small (16.2% for Moscosa perennials; 14.3% for Moscosa annuals; 11.9% for Urbión), but this was nevertheless the most important driver of spatially structured species composition in the Moscosa data. Spatial structure not correlated with hydrology ([b] + [c] + [e] in Fig.5), amounted to 13.4% for perennials and 12.4% for annuals. By comparison, species composition in the Urbión data was even more strongly spatially structured than in the Moscosa data, and more than a quarter of its spatial variance (26.7%) was not accounted for by hydrology. Nonetheless, we can conclude that hydrology was at least as important as any other single cause of spatial structure because it correlates with the primary axis of variation in Fig.1. Other causes of spatial structure could have included such ecological factors as vegetation pattern created by clonal growth, local variation in soil nutrients, or population processes such as dispersal (Legendre & Legendre 2012).

Our fifth question was whether annuals behaved differently from perennials and we found that indeed they did, occupying a distinct zone of niche space at the drier end of the hydrological gradient at Moscosa farm. Elsewhere, annuals have been found to partition a hydrological gradient in vernal pools in California (Bauder 2000), where even different genotypes of a single species occupy different zones of water depth (Linhart & Baker 1973). Niche segregation has also been experimentally demonstrated in several annual communities, though without always identifying the precise nature of the niche axes that are important for this (Sharitz & McCormick 1973; Turnbull, Manley & Rees 2005; Levine & HilleRisLambers 2009).

Our analysis has demonstrated that niche segregation along soil-moisture gradients contributes significantly to community structure, but the results are based in observational data and therefore offer only circumstantial evidence of the importance of the hydrological niche to
coexistence (Silvertown 2004). Complementary experimental and theoretical studies are required to quantify what contribution hydrological niche segregation makes to coexistence. We must also be cautious about how the soil-moisture gradient influences plant distribution, since soil moisture has direct and indirect effects upon the soil environment for plants. It not only controls water availability, but also when present in excess it affects oxygen availability, microbial community composition and function, and nutrient availability (Araya, Gowing & Dise 2012). That said, our dissection of the hydrological niche offers a firm statistical justification for exploring the underlying mechanisms and their consequences.
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Supporting Information

Additional supporting information may be found in the online version of this article.

Appendix S1. Sites photographs, full species names and species frequencies.

Appendix S2. Generalised Additive Mixed Modelling summaries.

Appendix S3. R code

Appendix S4. Moscosa dataset

Appendix S5. Lifespan
Table 1. Marginal tests results from direct RDAs (Redundancy Analysis) fitting groups of spatial MEM variables, linear trends, and the hydrological descriptor (AWTD) to explain species composition in a dehesa meadow at Moscosa Farm and in an alpine meadow at Urbión Peaks, Spain. The response is a Hellinger-transformed presence-absence species matrix in both cases. The hydrological component includes first-, second- and third-degree AWTD terms. The linear trend component is a surface described by the X-Y coordinates. The spatial component comprises sets of MEM spatial variables created specifically for each test (14 variables for the whole community and annuals in Moscosa; 15 for perennials in Moscosa; 18 for the whole community in Urbión); these MEM spatial variables describe spatial structuring. In complex models the amount of variation explained by each component depends on the other components (see Figure 5). \( p \)-values were obtained by means of 1000 permutations.

<table>
<thead>
<tr>
<th>Source of variation</th>
<th>Model var. (d.f.)</th>
<th>Resid var. (d.f.)</th>
<th>( F )</th>
<th>( p )</th>
<th>( R^2 ) (adj.)</th>
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</thead>
<tbody>
<tr>
<td>Whole community (Moscosa)</td>
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<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Hydrological descriptor</td>
<td>0.0806 (3)</td>
<td>0.3447 (192)</td>
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<tr>
<td>Linear trend</td>
<td>0.0631 (2)</td>
<td>0.3622 (193)</td>
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<td>Spatial MEM variables</td>
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<td>0.2856 (181)</td>
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<tr>
<td>Annuals (Moscosa)</td>
<td>Hydrological descriptor</td>
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<td>0.3560 (192)</td>
<td>15.1</td>
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<tr>
<td></td>
<td>Linear trend</td>
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<td>Spatial MEM variables</td>
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<th>0.3358 (192)</th>
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<table>
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<th>Whole community (Urbión)</th>
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**Figure 1.** Main species gradients for La Mina meadow in Moscosa Farm (a) and Laguna Larga meadow in the Urbión Peaks (b), as described by non–metric multidimensional scaling (NMDS), in multivariate space (Jaccard measure). Species scores are weighted averages scores. Species with relative frequency $\leq 0.05$ were not included. Moscosa annuals are indicated in red and perennials in green. Non-metric goodness-of-fit of the ordination: $R^2 = 0.967$ (Moscosa) and $R^2 = 0.989$ (Urbión). NMDS is a technique for unconstrained ordination, so the trend surface for AWTD in m (blue lines) was overlaid onto the species ordination only after the NMDS procedure was concluded. Significance of trend surface: $F = 24.3$, e.d.f. = 2.94, $p$-value < 0.000 (Moscosa) and $F = 34.1$, e.d.f. = 2.85, $p$-value < 0.000 (Urbión). In both cases, the hydrological gradient accounts for most of the variation observed in the first ordination axis (NMDS1); the levels of the contours depict the hydrological niches used in this work. Key to species: see Appendix 1

**Figure 2.** Fitted Generalised Additive Mixed Models (GAMM) for 16 perennial species (Moscosa). The smooth terms (smoothers) represent the estimated probability of presence (occurrence) with increasing Average Water Table Depth (AWTD in m) at La Mina site, Moscosa Farm, Western Spain. See Table S1 in Appendix S2, where significance ($F$- and $p$-values together with the estimated degrees of freedom) and adjusted $R^2$ are detailed. Note consistency with NMDS ordination

**Figure 3.** Fitted Generalised Additive Mixed Models (GAMM) for 16 annual species (Moscosa). The smoothers represent the probability of presence with increasing Average Water
Table Depth (AWTD in m) at the field site. See Table S2 in Appendix S2. Note consistency with NMDS ordination.

**Figure 4.** Fitted Generalised Additive Mixed Models (GAMM) for 16 species (Urbión). The smoothers represent the probability of presence with increasing Average Water Table Depth (AWTD in m) at the field site. See Table S3 in Appendix S2. Note consistency with NMDS ordination.

**Figure 5.** Venn diagrams showing how variation in the composition of the two plant communities was partitioned among a hydrological component, a linear trend component and a spatial component described by MEM variables. Numbers are adjusted $R^2$ values (%). The hydrological component includes AWTD and its second- and third-degree terms. The linear trend represents a surface described by the X-Y coordinates. The spatial component comprises sets of MEM spatial variables, selected for each partition specifically (14 MEM variables for the annuals community in Moscosa; 15 for the perennials community in Moscosa; 18 for the whole community in Urbión). The unique contributions of the hydrological, trend and spatial components are denoted by [a], [b] and [c], respectively. The fractions [f] and [g] correspond to spatially structured biological variation that is explained by the hydrological component, which is also spatially structured (induced spatial variation); fraction [f] is related to local topography at a fine scale; fraction [g] is related to local linear gradients (mainly local gradients of slope). Fraction [d] corresponds to linearly structured hydrological variation. Fraction [e] corresponds to linear variation that is shared by the spatial MEM variables (the MEM variables model both purely linear variation and any complex structures present in the data).
Fig. 1a

(a) NMDS ordination with AWTD
(b) NMDS ordination with AWTD
Fig. 2
Fig. 3

(a) Agrostis pourret

(b) Aira caryophylle

(c) Aphanes arvens

(d) Bellardia trixag.

(e) Brassica barnel

(f) Bromus hordeace

(g) Galium parisien

(h) Jasiona minlau

(i) Juncus bufonius

(j) Ornithopus perps

(k) Petrorhaga nart

(l) Silene gallica

(m) Trifolium glomere

(n) Trifolium striatu

(o) Trifolium subter

(p) Xolantha gutta