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Biodiversity patterns in a macroinvertebrate community of a temporary pond network

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Abstract. 1. Macroinvertebrate assemblages of temporary ponds are ideal model systems to explore biodiversity patterns and metacommunity ecology. In addition, the study of the environmental variables driving such biodiversity patterns is essential in establishing proper guidelines for the conservation of the singular fauna of temporary ponds, especially since such ponds are vulnerable systems.

2. We analysed the macroinvertebrate assemblages and environmental characteristics of 80 ponds spread across the Doñana National Park, SW Spain to (i) analyse macroinvertebrate β -diversity and metacommunity structure; and (ii) discern the main environmental and spatial drivers of these patterns.

3. The pond network was highly heterogeneous as temporary ponds were highly variable. Macroinvertebrate β -diversity partitioning showed that species replacement made the greatest contribution to total β -diversity while the contribution of nestedness was small. The macroinvertebrate community structure and β -diversity were similarly driven by: electrical conductivity (and co-variables alkalinity, pH, and ion concentrations), plant richness (and the co-variable pond surface area), maximum depth, marsh, and coastal proximity as well as two spatial descriptors extracted from Moran's eigenvector maps. The spatial descriptors indicated that large interpond distances were involved, suggesting that species dispersal limitations only take place over long distances in the area.

4. Those taxa that departed from the general nested pattern, termed idiosyncratic, significantly contributed to the maintenance of high pond network diversity through the species replacement and occurred within particular environmental conditions in the pond network.

5. These results reveal that environmental heterogeneity and connectivity are key factors in the preservation of high macroinvertebrate diversity in nested pond networks with high numbers of idiosyncratic species.

Key words. Aquatic insects, environmental heterogeneity, idiosyncratic species, Mediterranean temporary ponds, nestedness, pond conservation, spatial descriptors, species replacement, turnover.

Introduction

The metacommunity is an emergent concept that considers the impact of the exchange of species in heterogeneous environments (Leibold *et al.*, 2004; Urban & Skelly, 2006). Temporary ponds, which are characterised by annual inundation-desiccation cycles (Williams, 1997), are ideal model systems to study metacommunity ecology given their simple structure, local abundance, and occurrence in pond networks that demonstrate clear environmental gradients (Vanschoenwinkel *et al.*, 2007; Pandit *et al.*, 2009). Although temporary ponds are widely distributed worldwide (Williams *et al.*, 2001), their high biodiversity contrasts with their sensitivity and vulnerability to external perturbation, which has led to great interest in their con-

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servation over the last few years (Williams *et al.*, 2001; Zacharias *et al.*, 2007; Céréghino *et al.*, 2008). In addition, temporary ponds harbour singular flora and fauna that are often exclusive or infrequently found in permanent ponds (Collinson *et al.*, 1995; Williams, 1997; Céréghino *et al.*, 2008). In particular, their singular macroinvertebrate species can adjust their life cycles to the annual period of pond inundation (hydroperiod), re-starting community assembly after each year's initial inundation (Bazzanti *et al.*, 1996; Boix *et al.*, 2004; Florencio *et al.*, 2009).

In metacommunity ecology, β -diversity, which is the variation in species composition among sites in a geographical area (Legendre et al., 2005; but see e.g. Tuomisto, 2010; Anderson et al., 2011), is a key concept for understanding ecosystem functionality from a management and conservation perspective. In pond networks, environmental heterogeneity has been revealed as crucial in supporting high biodiversity (Urban, 2004; Jeffries, 2005) and also in driving patterns of nested biodiversity, in which species-poor sites contain subsets of species-rich sites, particularly in those systems with good conservation status (Hylander et al., 2005; Florencio et al., 2011). Hence, the study of those species that depart from the expectations of nested biodiversity patterns, which occur more or less frequently than would be predicted in a nested system (termed idiosyncratic), is currently receiving great interest in applied ecology (e.g. Florencio et al., 2011). To better understand the ecological processes maintaining high ecosystem diversity, β-diversity should be partitioned between (i) the β-diversity associated with non-random species loss in nested systems; and (ii) the β-diversity associated with true species replacement (Baselga, 2010). It is essential to disentangle the problem whether β -diversity is driven by species replacement or nestedness to make appropriate conservation decisions. If the former is the driver, it would prioritise the conservation of a large number of sites with variable richness and environmental conditions, while the latter would prioritise the conservation of the richest sites (Baselga, 2010).

One of the main debates in metacommunity ecology involves the relative importance of deterministic, nichebased process (e.g. environmental filters) versus stochastic ecological process (e.g. dispersal filters) in community assembly (Chase & Myers, 2011). Water chemistry and the physical characteristics of ponds each have an important influence on macroinvertebrate composition and abundance in wetlands (Wissinger, 1999; Williams, 2006). Conductivity is one of the most frequent chemical descriptors of macroinvertebrate communities (Garrido & Munilla, 2008; Waterkeyn et al., 2008). In particular, acidic water has negative effects on macroinvertebrate species diversity (Radke et al., 2003). Although nutrient concentrations have controversial effects, they usually negatively impact species occurrences at high levels (Declerck et al., 2005). Applying the theory of island biogeography (Mac-Arthur & Wilson, 1967) to lakes and ponds, high macroinvertebrate and plant species richness is harboured in large ponds (Friday, 1987; Nicolet et al., 2004). Interpond distances can also affect the incidence of species in particular pond assemblages as a result of species dispersal limitations (Briers & Biggs, 2005; Sanderson *et al.*, 2005).

We explored the main drivers of β -diversity and community structure in a macroinvertebrate metacommunity in a pond network of excellent conservation status. This is a highly dynamic system in which thousands of ponds fill and desiccate annually, with only a few ponds retaining water during the summer. The novelty of our study resides in the fact that we obtained comparable data on macroinvertebrates in 80 ponds distributed across an extensive area. We hypothesised that (i) there is high biodiversity in the macroinvertebrate metacommunity, with species replacement and nestedness being the main contributors to β -diversity; (ii) environmental variability is key in maintaining such high macroinvertebrate diversity in the pond network; and (iii) both random (i.e. dispersal) and deterministic processes (i.e. environment) are operating together in the macroinvertebrate assembly. To evaluate these hypotheses, we used data from 80 ponds, collected over a single season, to analyse (i) if β-diversity was mainly sustained by nestedness or by species replacement, and (ii) if spatial connectivity and environmental variability had an important influence on macroinvertebrate structure and β -diversity.

Methods

Study area

Doñana National Park (SW Spain) is one of the most pristine wetlands in Europe; it was included in the RAMSAR convention in 1982 and was later designated as a World Heritage Site by UNESCO in 1995. This area is located between the mouth of Guadalquivir River and the Atlantic Ocean. In the park, there is a clear geomorphological distinction between the ancient northern area and the southern area, locally known as 'Marismillas', which has a more recent marine origin (Siljeström et al., 1994). The three main types of landscapes are as follows: a sandy area with stabilised dunes, a mobile dune system, and an extensive marsh area (see Siljeström et al., 1994 for a detailed geomorphological description of the area). The climate is Mediterranean sub-humid, with mild winters, hot and dry summers, and heavy rains falling mainly in autumn or winter (mean annual rainfall = 544.6 (SD 211.3) mm with significant interannual variability, see Díaz-Paniagua et al., 2010).

This area contains a pond network that is comprised of more than 3000 water bodies in wet years and that is mainly composed of temporary ponds spanning a wide range of hydroperiods (Díaz-Paniagua *et al.*, 2010). These ponds are fed by annual rainfall and a shallow water table that rises above the surface after heavy autumn or winter rainfall and they generally dry out during summer (Díaz-Paniagua *et al.*, 2010). The ponds are heterogeneous in surface area, depth, and hydroperiod and are very abundant in the stabilised dunes and areas of contact between the three types of

landscapes (Díaz-Paniagua et al., 2010). In Doñana, there are only two large permanent (or semi-permanent) ponds, which only occasionally dry out after successive years of severe drought. There are also artificially deepened ponds (hereafter referred to as zacallones, the local name) that supply water for cattle and wild fauna during summer. They are present across the whole park but are the main water bodies present in the southern areas. In the contact area between the stable dunes and the freshwater marsh, there are ponds filled by the running water of intermittent streams that mainly flow towards the marsh after heavy rains (hereafter referred to as caños, the local name). This study included ponds that are representative of those in the study area and that were randomly selected across the entirety of the park: 46 temporary ponds, one of the two semi-permanent ponds, 27 zacallones, and 6 caños; we have grouped them according to their location in the northern or southern areas of the park (Fig. 1).

Macroinvertebrate sampling and taxon identification

We carried out a single survey of 80 ponds (late Marchmiddle June of 2007) spread across the whole of Doñana National Park (SW Spain) to analyse the environmental and spatial effects operating over the minimum time win-



Fig. 1. Locations of the 80 study ponds in Doñana National Park: 46 temporary ponds, which were mainly located in the northern part of the park, 27 *zacallones*, which were mainly located in the southern part of the park, 6 *caños*, and 1 semi-permanent pond are indicated.

dow in which all sites could be visited. We determined the presence or absence of macroinvertebrates using a dip net $(39 \times 21 \text{ cm}, 1 \text{ mm} \text{ mesh size})$ and netting across a stretch of water of approximately 1.5 m length in each sampling unit. In each pond, we sampled all different available microhabitats, based largely on differences in aquatic plant cover and depth (Heyer et al., 1994). As the efficiency of dip netting increases in small ponds (Heyer et al., 1994), we took more samples in larger ponds, which also typically contained a higher number of microhabitats, to achieve a comparable effort in detecting rare species (samples per pond ranged from 3 to 13). Sampling appropriateness was supported by a previous study in which similar results were obtained for sample-based rarefaction and raw data (see Florencio et al., 2011 for details). Most macroinvertebrates were identified in situ and then released again into the pond. Only unidentified individuals were preserved in 70% ethanol for later identification in the laboratory. We identified individuals to the species or genus level, except for Basomatophora, Diptera, Oligochaeta, and saldid bugs, which were identified to the family level (see Appendix 1 for the detailed taxonomic list). For those species for which we identified larvae and adults, we considered both stages separately in our analyses because of their different environmental requirements; they are thus referred to as taxa stages in our data.

Environmental variables in the extensive macroinvertebrate survey

To characterise the environmental gradients in Doñana ponds, we considered different groups of variables.

Environmental variables. In the field, we visually identified all the different plant taxa (species or genus level) per pond to estimate plant richness (Rplant). Maximum water depth (Max depth) was measured at the deepest point of the pond with a graduated pole. Pond surface area, the total number of ponds with an extension $>150 \text{ m}^2$ into a 200 m buffer area around each pond, and the total flooded surface area in a 200 m buffer area around each pond were extracted from a GIS-based map of ponds obtained from a hyperspectral image taken at a moment of high inundation of the area (see Gómez-Rodríguez et al., 2008 for details). We recorded in situ pH (near the bed using pH meter HI 991000, HANNA instruments, Portugal), dissolved oxygen concentration (near the bed using YSI 550A Handheld Dissolved Oxygen and Temperature System, YSI Incorporated, Yellow Springs, OH, USA), electrical conductivity (EC) at 20 °C (near the bed using Multi-range Conductivity Meter HI 9033, HANNA instruments, Romania), and turbidity (in the water column using Loggin Microprocessor turbidity meter HI 93703-11, HANNA instruments, Hungary). Surface water (500 ml) was collected to determine nutrient concentrations (dissolved inorganic phosphate, nitrate, nitrite, and ammonium), alkalinity, and main cation and anion concentrations (Cl^{-} , Na^{+} , Ca^{2+} ,

 K^+ and Mg^{2+}). Ion concentrations were analysed using a Trace Inductively Coupled Plasma Mass Spectrometer, while nutrient concentrations were measured colourimetrically using an Auto Analyser (Bran+Luebbe). Alkalinity was analysed according to the titration method described in APHA (1998). Surface sediment samples (5 cm depth) were collected and the following variables were measured in the laboratory: organic matter (three replicates; lost on ignition, 450 °C, 5 h) and sediment total P (two replicates). Sediment total P was estimated using dissolved inorganic phosphate obtained following the method of Murphy and Riley (1962), in which the ignited sediment undergoes acid digestion with 0.5 M H₂SO₄ and K₂S₂O₈ (0.5-1 g) at 120 °C for 4 h (Golterman, 2004). The total Fe concentration in digested sediment (two replicates) was determined colourimetrically by means of o-phenantroline and using ascorbic acid as the reducing agent (Golterman, 2004). The Na^{+}/Ca^{2+} ratio was measured because of its biological implications in regulating processes associated with the acid-base balance of the organisms (Radke et al., 2003). We did not use nitrite and nitrate concentrations in the analyses because most values were negligible (range <0.15- $0.60 \text{ mg } l^{-1}$).

Marsh-coast distance variables. To account for the influence of potential external sources of organisms (see e.g. Fahd *et al.*, 2007), we measured the minimum linear distances from each pond to the border of the marsh (Dmarsh) and the coast (Dcoast); these distances were also estimated using the GIS pond map (see Gómez-Rodríguez *et al.*, 2008 for details).

Spatial variables. Seventy-nine orthogonal spatial descriptors based on interpond distances were generated using Moran's eigenvector maps (MEMs) in R software 2.11.1 (R Development Core Team, 2010) ('spacemakeR' package, Dray, 2010; see Dray et al., 2006), which provide a general framework of principal coordinates of neighbour matrices (see Borcard & Legendre, 2002). The spatial descriptors extracted from the MEMs were ordered from V1 to V79, i.e. from the highest to the lowest eigenvalues. A selection of spatial descriptors that controlled for Type I error in the analyses was carried out according to Peres-Neto and Legendre (2010). The number of spatial descriptors was reduced using the 'ortho.AIC' command in R software 2.11.1 (R Development Core Team, 2010) ('spacemakeR' package, Dray, 2010). Only significant spatial descriptors with positive eigenvalues were considered in the analyses described below [redundancy analysis (RDA) and variation partitioning] to evaluate the effect of interpond distances on the structure of macroinvertebrate assemblages.

Statistical analyses

We constructed a pond-characteristic matrix with the values of the environmental and marsh-coast distance

variables. In addition, each group of variables (environmental, marsh-coast distance, and spatial) was considered in three individual matrices. Each variable had been previously transformed to approximate normality (Appendix 2). To obtain the pond-characteristic resemblance matrix, Euclidean distance was applied to the pond-characteristic matrix (Legendre & Legendre, 1998). Finally, we constructed a macroinvertebrate matrix that included the number of samples in which every taxa stage was present divided by the total number of samples taken in a pond. The Bray-Curtis index was applied to the macroinvertebrate matrix to obtain the macroinvertebrate resemblance matrix (Legendre & Legendre, 1998). Subsets of the macroinvertebrate matrix were extracted for the main taxonomical orders Coleoptera, Heteroptera, and Odonata.

To visualise the environmental variability in pond characteristics, we represented the pond dissimilarities by performing non-metric multidimensional scaling (NMDS) in PRIMER version 6 (Clarke & Warwick, 2001) on the pond-characteristic matrix.

We calculated the mean pair-wise macroinvertebrate β diversity (β sor) in our extensive sampling survey data to analyse macroinvertebrate β -diversity in the study area. The Sørensen index was applied to the presence-absence data (Legendre & Legendre, 1998). Using R software 2.11.1 (R Development Core Team, 2010), we partitioned β sor into β -diversity associated with species replacement (βsim) and β -diversity associated with nestedness (βnes) using the pair-wise measure approach described in Baselga (2010). In short, the total dissimilarity between each pair of ponds (β sor) was partitioned into two additive components accounting for dissimilarity due to species replacement (β sim) and dissimilarity due to nestedness (β nes), respectively, following the formula $\beta sor = \beta sim + \beta nes$ (Baselga, 2010). We also performed β-diversity partitioning using monthly macroinvertebrate assemblages of 22 of the temporary ponds for 2 years with different rainfalls (see Florencio et al., 2009, 2011 for a detailed description of macroinvertebrate sampling). As we obtained similar results, these data are not presented here for the sake of simplicity.

To detect which environmental variables influenced the macroinvertebrate assemblage structure of ponds, we performed constrained ordination using RDA in R software 2.11.1 (R Development Core Team, 2010) ('vegan' package, Oksanen et al., 2010) on each of the environmental, marsh-coast distance, and spatial variable matrices and the macroinvertebrate matrix and, independently, on the Coleoptera, Odonata and Heteroptera matrices. In the RDA, we excluded taxa stages that occurred in less than five ponds (30% of total species number) to avoid the disrupting effect of rare species (Leps & Smilauer, 2003). To exclude co-variables found to have poor explanatory power in RDA, we performed Spearman rank correlations (r_s) between each pair of environmental variables (Appendix 2). We used a forward stepwise procedure to select environmental variables, as described in Blanchet et al. (2008). Variation partitioning was performed in R soft-

ware 2.11.1 (R Development Core Team, 2010) ('vegan' package, Oksanen *et al.*, 2010) to measure the independent effects of environmental, marsh-coast distance, and spatial variables (see Borcard *et al.*, 1992); only explanatory variables found to be significant were extracted from RDA and included. In our variation partitioning, we used the adjusted multiple coefficient of determination (Adj. r^2), as required when matrices have different numbers of variables (Peres-Neto *et al.*, 2006). Significances were tested using Monte Carlo permutation tests (999 permutations).

After the RDA were performed, the influence of the significant explanatory variables on particular taxa stages and assemblages was analysed by performing a LINKTREE analysis in PRIMER version 6 (Clarke & Warwick, 2001) on the macroinvertebrate matrix (our parameterisation used three as the minimum group size and four as the minimum split size). SIMPROF analyses retained divisions significant at the 0.05 level and yielded a dendrogram of the results, otherwise known as a linkage tree (Clarke et al., 2008). The pair-wise differences between each group of macroinvertebrate assemblages detected by the linkage tree were assessed using one-way ANOSIM analysis (ANOSIM statistic R is close to one when groups are completely different). We determined which taxa stages contributed the most to these pair-wise differences (>10% of contribution) using one-way simper analysis in PRIMER version 6 (Clarke & Warwick, 2001).

To explore the relative influence of environmental variables on the partitioning of β -diversity, we used multiple regression on distance matrices (MRM), an extension of Mantel test (Legendre *et al.*, 1994). Spearman correlations (r_s) were used in the MRM analyses. Significant explanatory variables were identified using a forward-selection procedure (Legendre *et al.*, 1994). The significance of MRM models was assessed using 1000 permutations and only the most significant non-correlated variables were retained within each group of variables (spatial descriptors, environmental and marsh-coast distance variables). We constructed three successive models: (i) *the spatial model*, which used significant spatial descriptors to measure spatial influences on β -diversity; (ii) *the spatial/envi-*

ronmental model, which used significant environmental variables in addition to spatial descriptors to obtain partial effects; (iii) *the spatial/environmental/marsh-coast distance model*, which added significant marsh-coast distance variables to the previous model. All these calculations were performed using R software 2.11.1 (R Development Core Team, 2010; 'ecodist' package, Goslee & Urban, 2007).

Results

Variability in pond characteristics

The NMDS representation of the environmental variables of the sampled ponds revealed a heterogeneous pond network (Fig. 2). Northern ponds, which were mainly temporary, evidenced their high environmental variability when compared with southern ponds, which were mainly *zacallones* (Fig. 2).

Macroinvertebrate β -diversity partitioning

We recorded 135 taxa stages across the 80 study ponds, with an average of 23.5 (SD 8.5) taxa stages per pond. Using our extensive macroinvertebrate survey, we found that β -diversity was important in the study area [pond average of $\beta sor = 0.65$ (SD 0.11)]. In β -diversity partitioning, species replacement contributed more to β diversity [pond average of $\beta sim = 0.52$ (SD 0.14)] than nestedness [pond average of $\beta nes = 0.13$ (SD 0.10)].

Pond environmental variables influencing macroinvertebrate community structure

The significant explanatory variables influencing pond macroinvertebrate compositions detected by RDA were EC, Rplant, and Max depth among the environmental variables; both Dmarsh and Dcoast; and two spatial



Fig. 2. Non-metric multidimensional scaling ordination of the 80 study ponds according to the pond-characteristic resemblance matrix (Euclidean distance). Temporary ponds (Temp), *zacallones* (z), *caños* (Caño), and the semi-permanent pond (Semip) are highlighted as well as the location of ponds in southern (South) and northern (North) areas of the park.

Explanatory variables	Community	Coleoptera	Odonata	Heteroptera
Environmental	<i>F</i> -ratio = 3.347**	<i>F</i> -ratio = 3.23**	<i>F</i> -ratio = 4.45*	<i>F</i> -ratio = 4.85**
EC	0.34*	0.02*		0.25**
Rplant	0.18*		_	0.15**
Max depth	0.17*	0.02**		0.10*
Pond area		0.03**	n.s.	
Na^+/Ca^{2+} ratio		n.s.	n.s.	$\overline{0.08*}$
Pond number				0.08*
Alkalinity	_	_	$\overline{0.05}^{*}$	
Marsh-coast distance	F-ratio = 2.898**	F-ratio = 2.75**	<i>F</i> -ratio = 3.64*	
Dmarsh	0.26**	0.02**	0.04*	n.s.
Dcoast	0.17*	0.03**		n.s.
Spatial	<i>F</i> -ratio = 2.423**	<i>F</i> -ratio = 1.89*		n.s.
V5	0.17*	0.02*	n.s.	n.s.
V2	0.15*	n.s.	n.s.	n.s.

Table 1. Significant explanatory variables emerging from redundancy analyses (on the whole macroinvertebrate community and Coleoptera, Odonata, and Heteroptera matrices) performed independently on environmental, marsh-coast distance, and spatial variables[†].

n.s., non-significant variables; _, excluded variables; EC, electrical conductivity; Rplant, plant richness; Max depth, maximum water depth; Pond area, pond surface area; Pond number, total number of ponds with an extension > 150 m^2 into a 200 m buffer area around each pond; Dmarsh, minimum linear distances from each pond to the border of the marsh; Dcoast, minimum linear distances from each pond to the coast; V5, V2, eigenvectors extracted from the inter-pond distance based on the Moran's eigenvector maps.

[†]Values are the explained variance, indicating the magnitude of the effects of each significant explanatory variable, and global *F*-ratios. *P < 0.05.

**P < 0.01.

descriptors with high eigenvalues, V2 and V5. These high eigenvalues implied large interpond distances were involved (Table 1). EC had the greatest effect on the macroinvertebrate community (Table 1).

Coleopterans averaged 10.8 (SD 5.2) taxa stages per pond. We found that three groups of environmental variables had important effects on the structure of coleopteran assemblages. EC, Max depth, and Pond surface area were the significant environmental explanatory variables; both marsh-coast distance variables, Dmarsh and Dcoast, were significant; and only the spatial descriptor V5 had a significant effect among the spatial variables (Table 1). Odonatan assemblages [average = 2.4 (SD 2.1) taxa per pond] were significantly explained by Alkalinity and Dmarsh; no spatial descriptors were significant explanatory variables (all P > 0.36, Table 1). For heteropteran assemblages [average = 7.1 (SD 2.8) taxa stages per pond], EC, Rplant, Max depth, Na⁺/Ca²⁺ ratio, and the total number of ponds with an extension $>150 \text{ m}^2$ into a 200 m buffer area around each pond were significant environmental explanatory variables; no marsh-coast distance variables or spatial descriptors were significant explanatory variables (all P > 0.09, Table 1).

Variation partitioning analyses revealed that environmental variables (EC, Rplant and Max depth) had a more important effect on macroinvertebrate assemblage structures than marsh-coast distance and spatial variables (Table 2). Environmental variables were also the most important explanatory variables in coleopteran (EC, Max depth and Pond surface area), odonatan (Alkalinity), and heteropteran assemblages (EC, Rplant, Max depth, Na⁺/

Table 2. Independent effects of environmental, marsh-coast distance, and spatial variables on macroinvertebrate community structure and Coleoptera, Odonata, and Heteroptera matrices as indicated by variation partitioning analyses.

Adj. $r^{2\dagger}$	Community	Coleoptera	Odonata	Heteroptera
Environmen- tal	0.10**	0.07**	0.04**	0.20**
Marsh-coast distance	0.05**	0.03**	n.s.	_
Spatial	0.02**	0.01*	_	-

[†]Adjusted r^2 (ranged 0–1).

*P < 0.05.

**P < 0.01.

 Ca^{2+} ratio in water column, and the total number of ponds with an extension >150 m² into a 200 m buffer area around each pond) (Table 2). There were no significant independent effects of marsh-coast distance variables and spatial descriptors on the structure of odonatan and heteropteran assemblages (Table 2).

Pond macroinvertebrate assemblages and environmental thresholds

The linkage tree differentiated 16 pond groups based on differences in EC, Max depth, Dcoast, Dmarsh, Rplant, and pond macroinvertebrate assemblages. Macroinvertebrate assemblages associated with these pond



Fig. 3. Linkage tree representation showing significant divisive clustering of pond macroinvertebrate assemblages constrained by the significant environmental and marsh-coast distance variables detected by redundancy analyses: electrical conductivity (μ S cm⁻¹), maximum depth (cm), distance to the coast (Km^c), distance to the marsh (Km^m), and plant richness (Rplant). Pond number is indicated in each split group. Each successive split is conditioned by the indicated environmental thresholds of previous splits. B% is the contribution of each binary partition to global dissimilarity (ranged 0–100%). Capital letters indicate main taxa-stage contributors (>10% of contribution) to each split group (-A is adults, -L is larvae). R is the Spearman coefficient giving the dissimilarity value in every split.

groups differed in their contribution to the global dissimilarity of the whole macroinvertebrate community (Fig. 3). Fourteen taxa stages were the main contributors to pond assemblage differences along a generalist-specialist gradient; species ranged from occurring in several different types of environments to only being recorded under specialised conditions (Fig. 3). Four generalist taxa stages occurred under multiple environments (adults of Corixa affinis Leach, 1817; adults of Anisops sardeus Herrich-Schäffer, 1849; larvae of Cloeon Leach, 1815 spp.; and larvae of Notonectidae); another five taxa stages were favoured by narrower environmental conditions [Sympetrum fonscolombei (Selys, 1841) larvae; Chironomus plumosus (Linneo, 1758) larvae; Anacaena lutescens (Stephens, 1829) adults; Gerris thoracicus Schummel, 1832 adults; and Gerris spp. larvae]. Five further specialist taxa stages

were the main species contributing to the differentiation of ponds with particular characteristics: adults of Hydrobius fuscipes (Linnaeus, 1758) and Limnoxenus niger (Zschach, 1788) as well as Corixidae larvae and Culicidae larvae mainly occurred in shallow waters; Plea minutissima Leach, 1817 adults mainly occurred in deep ponds far from the coast with poor Rplant; and Sigara lateralis (Leach, 1817) adults were also common in deep waters with poor Rplant but that were close to the coast and far from the marsh (Fig. 3). Sympetrum fonscolombei larvae, Gerris spp. larvae, G. thoracicus adults, A. lutescens adults, H. fuscipes/L. niger adults, Corixidae larvae, C. plumosus larvae, Culicidae larvae, and P. minutissima adults made the greatest contributions to the global dissimilarity of the community, whilst S. lateralis adults contributed the least (Fig. 3).

Environmental variables driving the macroinvertebrate β -diversity pattern

Among the variables included in the MRM analyses, only NH₄ influenced the β *nes* pattern observed in the pond network ($r_s = 0.13$, $r^2 = 0.017$, P < 0.05). With

Table 3. Different multiple regression models associating macroinvertebrate β -diversity to the species replacement and including only spatial variables; spatial and environmental variables; and spatial, environmental, and marsh-coast distance variables.

Variable	$r_{\rm s}^{\dagger}$
Spatial model	$r^2 = 0.043^{**}$
V1 [‡]	0.153**
V5 [‡]	0.140**
Spatial/environmental model	$r^2 = 0.074^{**}$
V1	0.077 m.s.
V5	0.139**
Alkalinity	0.141**
Rplant	0.0949**
Max depth	0.069*
Spatial/environmental/marsh-coast distance model	$r^2 = 0.090^{**}$
V1	0.080*
V5	0.130**
Alk	0.136**
Rplant	0.086*
Max depth	0.063*
Dcoast	0.094*
Dmarsh	0.084*

m.s., marginally significant (P = 0.057); Rplant, plant richness; Max depth, maximum water depth; Dcoast, minimum linear distances from each pond to the coast; Dmarsh, minimum linear distances from each pond to the border of the marsh. [†]Coefficients of Spearman correlations, r^2 (ranged 0–1), 1000 permutations.

[‡]Eigenvectors extracted from the interpond distance based on the Moran's eigenvector maps.

*P < 0.05.

**P < 0.01

respect to the explanatory variables of βsim involved in β -diversity, two spatial descriptors with high eigenvalues (V1 and V5) were found to be significant variables in the spatial model (Table 3). In the spatial/environmental model, V5 and Alkalinity had the highest coefficients and thus best explained the βsim values (Table 3). In the spatial/environmental/marsh-coast distance model, V5 and Alkalinity were retained and shared similar, high coefficients that revealed their influence on βsim (Table 3). The spatial/environmental/marsh-coast distance model, which included the highest number of significant variables, explained 9% of variation in $\beta sim (r^2 = 0.09$, Table 3).

Discussion

Environmental variability

The high dissimilarity detected in the environmental characteristics of Doñana ponds reveals this system to be highly heterogeneous; this pattern is particularly due to the wide variability observed among temporary ponds, which are the most abundant aquatic habitats in this area. Although the artificially deepened ponds (zacallones) in the southern area of the park were more similar in their environmental characteristics, they widely differed from northern water bodies, thus increasing the heterogeneity of the total pond network. The long hydroperiod of these zacallones extends the temporal availability of aquatic habitats in the area and thus they act as reservoirs for species typical of temporary ponds, mainly macroinvertebrate dispersers that are forced to leave drying ponds in summer (see e.g. Garrido & Munilla, 2008; Florencio et al., 2009).

Macroinvertebrate β -diversity in a heterogeneous pond network

The macroinvertebrate β -diversity pattern reveals a diverse system mainly driven by species replacement in the pond network. Although the macroinvertebrate community of the Doñana pond network has been described as having a clear nested pattern (Florencio *et al.*, 2011), β -diversity partitioning indicated that nestedness hardly contributed to macroinvertebrate β -diversity. The relative importance of species replacement to β -diversity described in this study is concordant with the high number of idio-syncratic taxa stages (59) and ponds (34) found in the area that departed from the general nested pattern (Florencio *et al.*, 2011).

We detected some species whose occurrence was associated with particular environmental conditions, supporting the role of pond environmental heterogeneity in driving species replacement. In the linkage tree, we detected 10 taxa that were specialists occurring in a narrow range of environmental variability. These 10 specialists, with the exception of *Gerris* spp. larvae, were all included in the 59 idiosyncratic taxa stages listed for the Doñana pond network (see Florencio *et al.*, 2011). Except for *S. lateralis*, these taxa stages significantly contributed to the global dissimilarity of the whole macroinvertebrate community.

Relationships between macroinvertebrate assemblages and pond characteristics

We found similar explanatory variables influencing the macroinvertebrate community structure and the β-diversity associated with species replacement: EC (and alkalinity as a co-variable), maximum depth, aquatic plant richness, and distance from the ponds to the marsh and the coast, and two spatial descriptors. In metacommunity ecology, patterns of β -diversity are usually driven by biogeographical conditions (i.e. closer ponds should be more similar than more distant ponds as a result of species dispersal capabilities) as well as by environmental heterogeneity associated with complex processes (Leibold et al., 2004; Legendre et al., 2005). In this study, we found that one spatial descriptor as well as pond variability in alkalinity (and the co-variables EC, pH, and ion concentration) drove the macroinvertebrate β -diversity pattern via species replacement. Therefore, spatial and environmental filters are operating in community assembly via dispersal and species-sorting respectively (Patrick & Swam, 2011). These results are concordant with Chase and Myers (2011)'s predictions: β-diversity increases across spatial gradients in accordance with stochastic dispersal processes and β-diversity increases across environmental gradients in accordance with the niche-based theory.

In this study, EC (and co-variables) was correlated with the distance of ponds from the coast, revealing a gradual increase in water conductivity values from the north to the south of the park (82–8800 μ S cm⁻¹). The study ponds have no surface or groundwater connection to the sea though they have some oceanic influence due to airbone sea salt deposition and so the closer to the coast the higher the electrical conductivity. This conductivity gradient thus influences both the macroinvertebrate community structure and the β-diversity pattern. We also found that some species typically occurred under low conductivity conditions, for example A. lutescens occurred in waters with values lower than 225 μ S cm⁻¹. Similarly, in other temporary water systems, different species can occur across wide conductivity gradients (see e.g. Gutiérrez-Estrada & Bilton, 2010). The occurrence of different species can be favoured at different values of the conductivity gradient. For example we observed that Heteropterans, for example the corixid S. lateralis, preferred southern zacallones, which exhibited the highest conductivity in the study area. In contrast, Odonatans preferred northern temporary ponds with the lowest conductivity values; for example S. fonscolombei was observed almost exclusively in these ponds.

When exploring the influence of interpond distances on macroinvertebrate assemblage structure and macroinverte-

brate β -diversity, we only obtained spatial descriptors with high eigenvalues, a result that, in natural systems, signifies the involvement of broad spatial scales (Borcard & Legendre, 2002; Diniz-Filho & Bini, 2005; Griffith & Peres-Neto, 2006). Therefore, in this study, only the largest interpond distances had an effect on the macroinvertebrate assemblages and thus also on macroinvertebrate β -diversity, resulting in a system with high connectivity where species demonstrated weak dispersal limitations. The excellent dispersal abilities that usually characterise species of temporary ponds and let them cope with pond desiccation (Williams, 2006) largely contributed to the weak dispersal limitations in the study area. The Doñana pond network has already been determined to be a robust network for amphibian species, allowing them to encounter reproduction habitats even in extremely dry years (Fortuna et al., 2006); we confirm in this study that this assessment also applies to macroinvertebrate species.

We identified aquatic vegetation as an important variable structuring the macroinvertebrate community and its diversity, which is concordant with other studies carried out in temporary waters (e.g. Nicolet et al., 2004; Bilton et al., 2009). Diverse vegetation offers a wide range of niches for macroinvertebrate species, with a high number of refuges for species under predation and food availability for grazers. Plant species' architecture can influence biological processes, for example predator-prey interactions and the presence of oviposition sites. Hence, in this study, ponds with high aquatic plant richness harboured distinct macroinvertebrate assemblages that contributed significantly to macroinvertebrate diversity as a whole. In addition to aquatic vegetation, other biotic factors can affect the macroinvertebrate communities of temporary ponds; for instance, predators may have a seasonal effect, which could constitute an important focus for further research.

Implications for conservation

We demonstrate that both stochastic and deterministic ecological processes can operate together to assemble macroinvertebrates in a pond network. Stochastic processes such as dispersal only influenced the macroinvertebrate community and β -diversity at large spatial scales, which reveal the high connectivity of the system. Environmental variability was consequently key in maintaining high biodiversity in this system. The macroinvertebrate β diversity pattern was mainly driven by species replacement, with different species occurring in different environments; in contrast, the contribution of nestedness to β-diversity was low. Although the Doñana pond network has been described as having a clear nested pattern, the species that most contributed to β-diversity were largely idiosyncratic species and thus departed from the general nested pattern. We found that these idiosyncratic species occurred in specialised environments and were predominantly responsible for maintaining the system's high

biodiversity. In this study, we demonstrate the importance of idiosyncratic species in sustaining diversity in nested systems that contain high numbers of idiosyncratic species. Therefore, the best strategy for conservation is to preserve diverse environments across a non-fragmented habitat where species are not limited by dispersal. In other words, it is preferable to protect a wide range of diverse and interconnected ponds rather than the richest ones, which would be the conservation priority in a strictly nested system.

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Appendix 1. List showing the taxa captured in the study ponds (A is for adults, L is for larvae, N is for nymphs). The percent occurrence across all ponds (total), as well as in temporary ponds, zacallones, caños, and the semi-permanent pond is provided. Percentages for ponds located in the northern and southern areas of the park are given separately

Taxa		Total		Tempo	rary	Zacalle	sənes	Caños		Semi- perman	lent	Northe park	uria	Southe park	E
	Family	A	Γ	A	Γ	A	Γ	A	Γ	A	Γ	A	Г	A	Γ
Acari															
Hydrachnellae Bassomatonhora	I	3		0		7		0		0		7		9	
	Physidae	33		28		30		67		100		37		17	
Planorbidae	Planorbidae	14		6		19		17		100		<u>,</u> ∞		33	
Neotaenioglossa															
Potamopyrgus antipodarum	Hydrobiidae	3		0		7		0		0		0		11	
Coleoptera															
Bagous spp.	Curculionidae	16				7		0		100		21		0	
Dryops spp.	Dryopidae	34	1	39	2	30	0	0	0	100	0	40	7	11	0
Agabus conspersus (Marsham 1802)	Dytiscidae	39				59		17		100		34		56	
Agabus didymus (Olivier,	Dytiscidae	13		11		19		0		0		9		33	
1795)															
Agabus nebulosus (Forster, 1771)	Dytiscidae	21				22		33		100		24		11	
Agabus spp.	Dytiscidae		10		13		4		0		100		13		0
Cybister lateralimarginalis (De Geer, 1774)	Dytiscidae	б	24	0	26	4	19	0	17	0	100	0	24	9	22
Dytiscus circumflexus	Dytiscidae	4	6	4	11	4	٢	0	0	0	0	б	10	9	9
Fabricius, 1801		¢				c		¢		¢		,		¢	
<i>Eretes griseus</i> (Fabricius, 1781)	Dytiscidae	m		4		0		0		0		m		0	
Graptodytes flavipes (Olivier, 1795)	Dytiscidae	б		0		4		0		0		0		9	
Hydaticus leander (Rossi, 1790)	Dytiscidae		б		4		0		0		0		б		0
Hydroglyphus geminus (Fabricius, 1792)	Dytiscidae	33		24		48		17		100		23		67	
Hydroporus gyllenhali Schiödte. 1841	Dytiscidae	10		13		4		17		0		13		0	
Hydroporus lucasi Reiche,	Dytiscidae	45		46		48		17		100		45		44	
1000	Dytiscidae	26		13		56		0		0		11		78	

		Total		Temp	orary	Zacall	ones	Caños		Semi- permane	snt	Northe park	ern	Southe park	srn
Taxa	Family	V	Γ	A	Γ	A	Г	A	Г	A	Γ	A	Г	A	Γ
Hygrotus confluens (Fabricius, 1787)															
Hygrotus inaequalis (Fabricius, 1777)	Dytiscidae	б		0		Г		0		0		0		11	
Hygrotus lagari (Fery, 1992)	Dytiscidae	41		30		59		50		0		32		72	
<i>Hydroporus</i> spp. or <i>Hverotus</i> spp.	Dytiscidae		21		26		15		0		100		19		28
Hyphydrus aubei Ganolhaner 1802	Dytiscidae	20	20	13	13	33	37	17	0	0	0	13	15	44	39
Ilybius montanus (Scephens, 1828)/Agabus bipustulatus (1 innaeus 1767)	Dytiscidae	14		11		22		0		0		×		33	
<i>Laccophilus minutus</i> (Linnaeus, 1758)	Dytiscidae	35	51	33	50	41	56	33	33	0	100	34	50	39	56
Liopterus atriceps (Sharp, 1882)	Dytiscidae	10		11		4		33		0		13		0	
Rhantus hispanicus Sharp, 1882	Dytiscidae	36		39		30		33		100		44		11	
Rhantus suturalis (McLeay, 1825)	Dytiscidae	16		15		22		0		0		15		22	
Colymbetes fuscus (Linnaeus, 1758)	Dytiscidae	41		39		52		17		0		37		56	
Rhantus spp. or Colymbetes fuscus	Dytiscidae		20		30		4		0		100		26		0
Yola bicarinata (Latreille, 1804)	Dytiscidae	9		0		19		0		0		0		28	
Gyrinus dejeani Brullé, 1832	Gyrinidae	19		15		30		0		0		11		44	
Gyrimus urinator Illiger, 1807 Gyrinus sob.	Gyrinidae Gvrinidae	m	б	0	4	4	0	0	0	0	0	0	ŝ	9	0
Haliplus andalusicus	Haliplidae	9		7		7		0		0		ю		17	
Wehncke, 1874						;								:	
Haliplus guttatus Aubé, 1836 Haliplus lineatocollis (Marcham, 1807)	Haliplidae Haliplidae	s ss		0 0		==		0 0		0 0		n n		11	
Haliplus spp.	Haliplidae		9		6		4		0		0		9		9
Helophorus spp.	Helophoridae	45		46		41		50		100		48		33	
Limnebius furcatus Baudi, 1872	Hydraenidae	-		0		4		0		0		0		9	
Ochthebius dilatatus Stephens, 1829	Hydraenidae	\mathfrak{c}		4		0		0		0		б		0	

Appendix 1. (Continued).

		Total		Temp	orary	Zacal	lones	Caños		Semi- permar	hent	North park	ern	Southe park	E
Taxa	Family	A	Γ	V	Γ	V	Γ	V	Γ	A	Г	A	Γ	V	Γ
Ochthebius auropallens Fairmaire 1870	Hydraenidae	5		٢		4		0		0		3		11	
Hydrochus flavipennis Küster 1852	Hydrochidae	4		0		7		0		0		3		9	
Anacaena lutescens (Stephens, 1829)	Hydrophilidae	54		61		37		67		100		60		33	
Berosus affinis Brullé, 1835	Hydrophilidae	21		22		22		17		0		19		28	
Berosus guttalis Rey, 1883	Hydrophilidae	15		17		11		17		0		18		9	
Berosus signaticollis (Charnentier 1825)	Hydrophilidae	16		24		٢		0		0		19		9	
Berosus snn.	Hvdronhilidae		26		35		11		17		100		32		9
Enochrus bicolor (Fabricius, 1792)	Hydrophilidae	23	1	22		26		0	•	100	0	21		28)
Enochrus fuscipennis (C.G. Thomsom, 1884)	Hydrophilidae	40		46		33		33		0		45		22	
Enochrus spp.	Hydrophilidae		8		13		0		0		0		10		0
<i>Helochares lividus</i> (Forster, 1771)	Hydrophilidae	11		6		11		33		0		8		22	
Hydrobius convexus Brullé, 1835	Hydrophilidae	÷		÷											
Hydrobius fuscipes	Hydrophilidae	41		52		19		50		100		48		17	
(Linnaeus, 1758) & Linnoxenus niger (Zschach,															
1/88)									,						,
Hydrobius spp. or Limnoxenus niger	Hydrophilidae		28		37		19		0		0		35		0
Hydrochara flavipes (Steven, 1808)	Hydrophilidae	S	б	4	4	0	0	17	0	100	0	9	3	0	0
Hydrophilus pistaceus (Laporte, 1840)	Hydrophilidae		4		0		٢		17		0		5		0
Laccobius revelierei Perris, 1864	Hydrophilidae	-		7		0		0		0		7		0	
Paracymus scutellaris (Rosenhauer, 1856)	Hydrophilidae	20		24		11		33		0		26		0	
Hygrobia hermanni (Fabricius, 1775)	Paelobiidae	34	19	24	24	56	15	17	0	0	0	23	21	72	11
Noterus laevis Sturm, 1834	Noteridae	8		4		7		17		100		8		9	
<i>Hydrocyphon</i> spp. Ephemeroptera	Scirtidae		б		4		0		0		0		ŝ		0
Cloeon spp.	Baetidae		74		76		81		17		100		69		89

Appendix 1. (Continued).

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(Continued).
÷
Appendix

		Total		Tempo	rary	Zacallo	nes	Caños		Semi- permane	nt	Norther park	Ħ	Souther park	Ħ
Taxa	Family	A	Г	A	Г	A	Г	A	Г	A	Г	A	Г	A	Γ
Caenis spp.	Caenidae		1		0		0		17		0		2		0
Tubificidae	Tubificidae	1		2		0		0		0		2		0	
Heteroptera															
Corixa affinis Leach, 1817	Corixidae	81		78		93		50		100		LL		94	
<i>Micronecta scholzi</i> (Fieber, 1860)	Corixidae	~		0		19		0		0		0		28	
Sigara lateralis (Leach, 1817)	Corixidae	45		39		63		17		0		35		78	
Sigara scripta (Rambur, 1840)	Corixidae	14		6		26		0		0		9		39	
Sigara selecta (Fieber, 1848)	Corixidae	ю		2		4		0		0		7		9	
Sigara stagnalis (Leach, 1817)	Corixidae	14		6		26		0		0		9		39	
Trichocorixa verticalis (Fieber, 1851)	Corixidae	15		17		15		0		0		13		22	
Corixidae spp.	Corixidae		56		59		59		17		100		53		67
Gerris cf maculatus Tamanini, 1946	Gerridae	10		11		7		17		0		11		9	
Gerris thoracicus Schummel, 1832	Gerridae	65		72		52		67		100		69		50	
Gerris spp.	Gerridae		59		74		33		50		100		69		22
<i>Microvelia pygmaea</i> (Dufour, 1833)	Microveliidae	4		٢		0		0		0		5		0	
Naucoris maculatus Fabricius, 1798	Naucoridae	6	10	4	6	19	15	0	0	0	0	9	9	17	22
Nepa cinerea Linnaeus, 1798	Nepidae	10	10	6	4	7	15	33	33	0	0	11	8	9	17
Anisops sardeus Herrich-Schäffer, 1849	Notonectidae	81		78		93		50		100		79		89	
Notonecta glauca Linnaeus, 1758 ssp. glauca	Notonectidae	23		28		15		0		100		24		17	
Notonecta glauca Linnaeus, 1758 ssp. meridionalis Poisson, 1926	Notonectidae	29		28		33		17		0		29		28	
Notonecta maculata Fabricius, 1794	Notonectidae	14		13		19		0		0		13		17	
Notonecta viridis Delcourt, 1909	Notonectidae	26		20		41		17		0		26		28	
Notonectidae spp.	Notonectidae Pleidae	33	75 10	24	83 11	52	70 11	0	33 0	100	100 0	29	77 8	44	67 17

Appendix 1. (Continued).								
		Total	Temporary	Zacallones	Caños	Semi- permanent	Northern park	Southern park
Taxa	Family	A L	A L	A L	A L	A L	A L	A L
Plea minutissima Leach, 1817								
Notostraca								
Triops mauritanicus (Ghigi, 1921)	Triopsidae	б	4	0	0	0	С	0
Spinicaudata								
Cyzicus grubei Simon, 1886	Cyzicidae	1	2	0	0	0	0	9
<i>Maghrebestheria maroccana</i> Thiéry, 1988	Leptestheriidae	ю	4	0	0	0	б	0
Anostraca								
Branchipus cortesi Alonso y Janme, 1991	Branchipodidae	1	5	0	0	0	2	0
Branchipus schafferi Fischer de Waldheim, 1834	Branchipodidae	1	7	0	0	0	7	0
Tanymastix stagnalis (Linnaeus, 1758)	Tanymastigiidae	1	7	0	0	0	2	0
Streptocephalus torvicornis (Waga, 1842)	Chirocephalidae	б	4	0	0	0	2	6
Odonata								
Aeshna affinis* Vander Linden, 1823	Aeshnidae	8	11	4	0	0	∞	9
Aeshna mixta Latreille, 1805	Aeshnidae	8	4	11	17	0	8	9
Coenagrion scitulum (Rambur. 1842)	Coenagrionidae	9	L	7	0	0	8	0
Ischnura elegans* (Vander Linden, 1820)	Coenagrionidae	28	30	26	0	100	31	17
Ischnura pumilio* (Charp., 1825)	Coenagrionidae	38	41	30	33	100	44	17
Lestes barbarus (Fabr.,	Lestidae	13	15	11	0	0	16	0
1798)								
Lestes dryas Kirby, 1890 Lestes macrostigma (Evieren 1836)	Lestidae Lestidae	10	11 2	11 0	0 0	0 0	11	0
Lestes virens (Charpentier, 1875)	Lestidae	6	7	22	0	0	3	28
Crocothemis erythraea (Brullé 1832)	Libellulidae	4	4	4	0	0	3	9
	Libellulidae	51	67	26	33	100	63	11

Macroinvertebrate β *-diversity in ponds* 19

		Total		Temp	orary	Zacal	lones	Caños		Semi- permaı	hent	North park	ern	Southe	Е
Taxa	Family	А	Г	A	Г	A	Γ	A	Г	A	Г	A	Г	A	Γ
Sympetrum fonscolombei (Selvs, 1841)															
Sympetrum meridionale (Selvs, 1841)	Libellulidae		19		33		0		0		0		24		0
Sympetrum sanguineum (Müller, 1764)	Libellulidae		24		37		7		0		0		29		9
Sympetrum striolatum (Charpentier. 1840)	Libellulidae		26		37		٢		33		0		34		0
Orthetrum brunneum* (Fonscolombe, 1837)	Libellulidae		1		0		4		0		0		0		9
Orthetrum cancellatum (Linneo. 1758)	Libellulidae		1		0		4		0		0		0		9
Orthetrum nitidinerve* (Selvs. 1841)	Libellulidae		1		0		4		0		0		7		0
		Γ	Z	Γ	Z	Γ	Z	Γ	Z	L	Z	L	Z	L	z
Diptera															
Chaoborus spp.	Chaoboridae	9		11		0		0		0		8			
Chironomus plumosus (Linneo, 1758)	Chironomidae	09		57		70		33		100		60		61	
Culicidae	Culicidae	29	16	43	22	7	11	17	0	0	0	35	18	9	11
Dixa spp.	Dixidae	5		6		0		0		0		9		0	
Dolichopodidae	Dolichopodidae	1		0		0		0		0		0			
Ephydridae	Ephydridae		б		0		4		0		0		0		11
Orthocladiinae	Chironomidae	11		11		11		17		0		11		11	
Rhagionidae	Rhagionidae	5		6		0		0		0		9		0	
Sciomyzidae	Sciomyzidae	1		0		0		0		0		0		0	
Tabanidae	Tabanidae	б		4		0		0		0		ς		0	
Tanypodinae	Chironomidae	6		7		7		33		0		8		11	
Tipulidae	Tipulidae	5		6		0		0		0		9		0	
		-			J. J	-				_	71		1 1		
[*] Confirmation of the identity	of these species is red	intred be	scause s	nectes 1d	entificat	ton kev	s based c	n larval	morph	ology are	C1111CI 1	to use an	nd these	species	lave

â 2, 5 Σ, not been previously cited as being in the study area. *Only occasional presence of this species was detected.

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Appendix 1. (Continued).

EnvironmentalMeanSDMeanSDNumberRplant7.43.73.93.014Rpant7.43.73.93.014Max depth (cm)59.831.8103.813.2118Pond area (m ²)339055861193291226725Pond number5.14.03.13.411Flooded area (m ²)331462681649339763<28212NH4 $^+$ (mg 1 ⁻¹)0.170.540.120.150.050.05NH4 $^+$ (mg 1 ⁻¹)0.110.130.080.100.01TP-s (µg g^{-1} d.w.)258.36314.59105.29121.80162.563Do 2 (mg 1 ⁻¹)0.110.130.080.100.017.61O2 (mg 1 ⁻¹)0.701.82.53.62.81.47D< 2 (mg 1 ⁻¹)0.71.82.362.617.61D< 2 (mg 1 ⁻¹)0.701.82.553.62.8D< 2 (mg 1 ⁻¹)0.724.964.221.47D< 2 (mg 1 ⁻¹)1.992.774.964.221.47D< 2 (mg 1 ⁻¹)0.726.602.633.493.93Malinity (meq 1 ⁻¹)275117183.42Na ⁺ (meq 1 ⁻¹)0.160.320.330.380.03Mg ²⁺ (meq 1 ⁻¹)0.160.320.330.380.03Mg ²⁺ (meq 1 ⁻¹)0.92 <th>$(N = 2^{\circ})$</th> <th>nes 7)</th> <th>Samin</th> <th>Caños (N</th> <th>= (9)</th> <th></th> <th></th>	$(N = 2^{\circ})$	nes 7)	Samin	Caños (N	= (9)		
Rplant 7.4 3.7 3.9 3.0 14 Max depth (cm) 59.8 31.8 103.8 13.2 118 Pond area (m ²) 3390 5586 119 329 122672 5 Pond number 5.1 4.0 3.1 3.1 3.1 3.1 3.1 Flooded area (m ²) 3390 5586 119 3.29 122672 5 NH4 ⁺ (mg 1 ⁻¹) 0.17 0.54 0.12 0.15 0.05 0.05 NH4 ⁺ (mg 1 ⁻¹) 0.11 0.13 0.54 0.12 0.15 0.05 0.05 NH4 ⁺ (mg 1 ⁻¹) 0.11 0.13 0.05 0.10 0.01 0.05 <td< th=""><th>SD Mean</th><th>SD</th><th>(N = 1)</th><th>Mean</th><th>SD</th><th>Transf.</th><th>$r_{ m s} > 0.6^{\dagger}$</th></td<>	SD Mean	SD	(N = 1)	Mean	SD	Transf.	$r_{ m s} > 0.6^{\dagger}$
$\begin{array}{llllllllllllllllllllllllllllllllllll$	3.7 3.9	3.0	14	6.2	3.5		Pond area (+)
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	31.8 103.8	13.2	118	52.2	34.2		
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	5586 119	329	122672	538	408	Log	Ag plant R (+)
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	4.0 3.1	3.4	11	2.8	2.2	, 1	Flooded area (+)
$\begin{array}{llllllllllllllllllllllllllllllllllll$	6268 1649	3397	63 282	1227	1993	Log	Pond number (+)
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	0.54 0.12	0.15	0.05	0.14	0.12	Log	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	0.13 0.08	0.10	0.01	0.21	0.28	Log	
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	314.59 105.25	121.80	162.56	328.40	333.52	I	TFe-s (+), O.M. (+)
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	3.18 2.36	2.61	7.61	4.57	4.41	I	TP-s (+), O.M. (+)
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	1.8 2.5	3.6	2.8	3.6	3.1	Log	
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	0.7 7.7	1.1	6.9	6.7	0.6	Log	EC (+), Alk (+), Na^+ (+), Ca^{2+} (+), K^+ (+), Mg^{2+} (+)
$ \begin{array}{rrrr} Alkalinity (meq \ l^{-1}) & 1.99 & 2.72 & 4.96 & 4.22 & 1.47 \\ Turbidity (NTU) & 27 & 51 & 17 & 18 & 3 & 1 \\ 0.M. (\%) & 7.02 & 6.60 & 2.63 & 3.49 & 3.93 \\ Cl^{-1} (meq \ l^{-1}) & 4.20 & 7.81 & 8.65 & 11.83 & 4.24 \\ Na^{+} (meq \ l^{-1}) & 3.88 & 6.47 & 7.28 & 7.87 & 4.3 \\ Ca^{2^{+}} (meq \ l^{-1}) & 0.85 & 1.15 & 2.19 & 2.41 & 0.59 \\ K^{+} (meq \ l^{-1}) & 0.16 & 0.32 & 0.33 & 0.38 & 0.03 \\ Mg^{2^{+}} (meq \ l^{-1}) & 0.92 & 1.34 & 3.34 & 3.86 & 1.05 \\ Na^{+}/Ca^{2^{+}} ratio & 14.49 & 58.08 & 4.92 & 4.80 & 7.24 \\ \end{array} $	1270.6 1304.7	1226.4	1224	479.2	356.3	Log	pH (+), Alk (+), CI^{-1} (+), Na^{+} (+), Ca^{2+} (+), K^{+} (+), Mg^{2+} (+), coast (-)
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	2.72 4.96	4.22	1.47	1.57	0.82	Log	pH (+), EC (+), CI^{-1} (+), Na^{+} (+), Ca^{2+} (+), K^{+} (+), Mg^{2+} (+)
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	51 17	18	б	192	385	Log	
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	6.60 2.63	3.49	3.93	6.08	6.00	I	TP-s (+) and TFe-s (+)
$ \begin{array}{llllllllllllllllllllllllllllllllllll$	7.81 8.65	11.83	4.24	3.79	2.91	Log	EC (+), Alk (+), Na^+ (+), Ca^{2+} (+), K^+ (+), Mg^{2+} (+)
$ \begin{array}{cccc} Ca^{2+} (meq \ l^{-1}) & 0.85 & 1.15 & 2.19 & 2.41 & 0.59 \\ K^{+} (meq \ l^{-1}) & 0.16 & 0.32 & 0.33 & 0.38 & 0.03 \\ Mg^{2+} (meq \ l^{-1}) & 0.92 & 1.34 & 3.34 & 3.86 & 1.05 \\ Na^{+}/Ca^{2+} ratio & 14.49 & 58.08 & 4.92 & 4.80 & 7.24 \\ \end{array} $	6.47 7.28	7.87	4.3	3.19	2.41	Log	EC (+), Alk (+), Cl ⁻¹ (+), Ca ²⁺ (+), K ⁺ (+), Mg ²⁺ (+)
$ \begin{array}{ccccc} K^{+} \mbox{ (meq } l^{-1}) & 0.16 & 0.32 & 0.33 & 0.38 & 0.03 \\ Mg^{2^{+}} \mbox{ (meq } l^{-1}) & 0.92 & 1.34 & 3.34 & 3.86 & 1.05 \\ Na^{+}/Ca^{2^{+}} \mbox{ ratio} & 14.49 & 58.08 & 4.92 & 4.80 & 7.24 \\ \end{array} $	1.15 2.19	2.41	0.59	0.67	0.47	Log	EC (+), Alk (+), pH (+), Cl ⁻¹ (+), Na ⁺ (+), K ⁺ (+), Mg ²⁺ (+)
$ \begin{array}{ccccc} Mg^{2+} (meq \ l^{-1}) & 0.92 & 1.34 & 3.34 & 3.86 & 1.05 \\ Na^{+}/Ca^{2+} \ ratio & 14.49 & 58.08 & 4.92 & 4.80 & 7.24 \\ \end{array} $	0.32 0.33	0.38	0.03	0.10	0.06	Log	EC (+), Alk (+), pH (+), CI^{-1} (+), Na^{+} (+), Ca^{2+} (+), Mg^{2+} (+)
Na^{+}/Ca^{2+} ratio 14.49 58.08 4.92 4.80 7.24	1.34 3.34	. 3.86	1.05	0.71	0.63	Log	EC (+), Alk (+), pH (+), CI^{-1} (+), Na^{+} (+), Ca^{2+} (+), K^{+} (+), Mg^{2+} (+)
	58.08 4.92	4.80	7.24	4.93	1.24	Log	
Marsh-coast distance							
Dmarsh (m) 2293 1798 2180 1450 3519 2	1798 2180	1450	3519	234	410	I	
Dcoast (m) 5097 2529 3085 2099 2168 95	2529 3085	2099	2168	9523	2806		EC (-)

coast distance variables) of sampled nonds march pue lotue otarietioe nond chara the for shown 040 deviation (SD) ž

streams which mainly flow towards the marsh after heavy rains (N = number of ponds); Transf., independent transformations to approximate normality (Sqr is Temporary, temporary ponds; Zacallones, artificially deepened ponds; Semip, the semi-permanent pond; Caños, ponds filled by the running water of intermittent square root transformation; Log is Log(X + 1) transformation; _, is no transformation required); Rplant, plant richness; Max depth, maximum water depth; Pond 02, dissolved oxygen concentration; EC, electrical conductivity; O.M., organic matter; Dmarsh, the minimum linear distances from each pond to the border of area, pond surface area; Pond number, total number of ponds with an extension >150 m² into a 200 m buffer area around each pond; Flooded area, total flooded surface area in a 200 m buffer area around each pond; i-P, dissolved inorganic phosphate; TP-s, sediment total P; TFe-s, total Fe concentration in the sediment; the marsh; Dcoast, the minimum linear distances from each pond to the coast.

Significant Spearman coefficient higher than 0.6 (all P < 0.01) indicating variables with positive (+) and negative (-) correlations.

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